

SEQUENCE LISTING

<110> Gudas, Jean M.
Haak-Frendscho, Mary
Foord, Orit
Liang, Meina L.
Ahluwalia, Kiran
Bhakta, Sunil

<120> ANTIBODIES DIRECTED TO MONOCYTE
CHEMO-ATTRACTANT PROTEIN-1 (MCP-1) AND USES THEREOF

<130> ABGENIX.091A

<150> 60/404,802
<151> 2002-08-19

<160> 149

<170> FastSEQ for Windows Version 4.0

<210> 1
<211> 1335
<212> DNA
<213> Homosapien

<400> 1
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tcctgcagg tttccggata caccctact gaattatcca tgcactgggt gcgacaggct 120
cctggaaatg ggcttgagtg gatgggaggt tttgatcctg aagatggta gacaatctac 180
gcacagagg tccaggggcag agtcgtcatg accgaggacc catctacaga cacagcctac 240
atggagctga gcaggcttag atctgaggac acggccgtgt attactgtgc aaccaacgag 300
ttttggagtg gttatattga ctactgggc cagggaaaccc tggtcaccgt ctccctcagcc 360
tccaccaagg gcccattcggt ctccccctg gcgcctgtct ccaggagcac ctccgagagc 420
acagcggccc tgggtgcct ggtcaaggac tacttccccg aaccgggtgac ggtgtcgtgg 480
aactcaggcg ctctgaccag cggcgtgcac accttcccgat ctgtcctaca gtccctcagga 540
ctctactccc tcagcagcgt ggtgaccgtg ccctccagca acttcggcac ccagacccatc 600
acctgcacg tagatcacaa gcccagcaac accaagggtgg acaagacagt tgagcgcaaa 660
tgttgtgtcg agtgcaccgc tgccccagca ccacctgtgg caggaccgtc agtcttcctc 720
ttccccccaa aacccaaggaa caccctcatg atctcccgaa cccctgaggt cactgtcgtg 780
gtggtggacg tgagccacga agaccccgag gtccagttca actggtaacgt ggacggcgtg 840
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gtcagcgtcc tcaccgttgt gcaccaggac tggctgaacg gcaaggagta caagtgcac 960
gtctccaaca aaggccccc agccccatc gagaaaaacca tctccaaaac caaagggcag 1020
ccccgagaac cacaggtgtc caccctgccc ccatcccggg aggagatgac caagaaccag 1080
gtcagcctga cctgcctgtt ccaaaggcttc taccccgacg acatcgccgt ggagtggag 1140
agcaatgggc agccggagaa caactacaag accacaccc ccatgctgga ctccgacggc 1200
tccttcttcc tctacagcaa gtcaccgtg gacaagagca ggtggcagca ggggaacgtc 1260
ttctcatgtt ccgtgatgca tgaggctctg cacaaccact acacgcagaa gagcctctcc 1320
ctgtctccgg gtaaa 1335

<210> 2
<211> 445

<212> PRT

<213> Homosapien

<400> 2

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Asn Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Arg Phe
50 55 60
Gln Gly Arg Val Val Met Thr Glu Asp Pro Ser Thr Asp Thr Ala Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr Asn Glu Phe Trp Ser Gly Tyr Phe Asp Tyr Trp Gly Gln Gly
100 105 110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125
Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
130 135 140
Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
145 150 155 160
Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu
165 170 175
Gln Ser Ser Gly Leu Tyr Ser Leu Ser Val Val Thr Val Pro Ser
180 185 190
Ser Asn Phe Gly Thr Gln Thr Tyr Thr Cys Asn Val Asp His Lys Pro
195 200 205
Ser Asn Thr Lys Val Asp Lys Thr Val Glu Arg Lys Cys Cys Val Glu
210 215 220
Cys Pro Pro Cys Pro Ala Pro Pro Val Ala Gly Pro Ser Val Phe Leu
225 230 235 240
Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu
245 250 255
Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Gln
260 265 270
Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys
275 280 285
Pro Arg Glu Glu Gln Phe Asn Ser Thr Phe Arg Val Val Ser Val Leu
290 295 300
Thr Val Val His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys
305 310 315 320
Val Ser Asn Lys Gly Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys
325 330 335
Thr Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser
340 345 350
Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys
355 360 365
Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln
370 375 380
Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Met Leu Asp Ser Asp Gly
385 390 395 400
Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln

405	410	415
Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn		
420	425	430
His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys		
435	440	445

<210> 3
<211> 660
<212> DNA
<213> Homosapien

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atcaactgta agtccagcca gagtgttta tacagctcca acaataagaa ctacttagtt 120
tggtaccagc agaaaaccagg acagcctcct aaactgctca tttactggc atctatccgg 180
gaatccgggg tccctgaccg attcagttcc agccgggtctg agacagattt cactctcacc 240
atcagcagcc tgcaggctga agatgtggca gtttattact gtcagcaata ttttagtagt 300
ccgtggacgt tcggccaagg gaccaaggtg gaaatcaaac gaactgtggc tgaccatct 360
gtcttcatct tcccgcacatc tgatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
ctgctgaata acttctatcc cagagagcc aaagtacagt ggaagggtgga taacgccctc 480
caatcgggta actcccagga gagtgtcaca gagcaggaca gcaaggacag cacctacagc 540
ctcagcagca ccctgacgct gagcaaagca gactacgaga aacacaaagt ctacgcctgc 600
gaagtcaccc atcagggcct gagctcggcc gtacacaaaga gcttcaacag gggagagtgt 660

<210> 4
<211> 220
<212> PRT
<213> Homosapien

<400> 4
Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Met Ser Leu Gly
1 5 10 15
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
20 25 30
Ser Asn Asn Lys Asn Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln
35 40 45
Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Ile Arg Glu Ser Gly Val
50 55 60
Pro Asp Arg Phe Ser Ser Gly Ser Glu Thr Asp Phe Thr Leu Thr
65 70 75 80
Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
85 90 95
Tyr Phe Ser Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
100 105 110
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
115 120 125
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
130 135 140
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu
145 150 155 160
Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp
165 170 175
Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr

180	185	190
Glu Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser		
195	200	205
Ser Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys		
210	215	220

<210> 5
<211> 475
<212> DNA
<213> Homosapien

<400> 5
caggtccagc tggcacagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcagg tttccggata caccctact gaattatcca tgcactgggt ggcacaggct 120
cctggaaaag ggcttgagtg gatggggaggt tttgatcctg aagatggtga aacaatctac 180
gcacagaagt tccaggcag agtcaccatg accgaggaca catctacaga cacagcctac 240
atggagctga gcagcctgag atctgaggac acggccgtgt attattgtgc aaccaacgaa 300
ttttggagtg gttatgttga ctactgggc cagggAACCC tggtcaccgt ctccctcagcc 360
tccaccaagg gcccattcggt ctccccctg gcgcctgtgt ccaggagcac tacttcccc 420
ggcgtgcaca cttcccccagc tgcctacag tcctcaggac tctactccct cagca 475

<210> 6
<211> 158
<212> PRT
<213> Homosapien

<400> 6
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr Asn Glu Phe Trp Ser Gly Tyr Phe Asp Tyr Trp Gly Gln Gly
100 105 110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125
Pro Leu Ala Pro Cys Ser Arg Ser Thr Thr Ser Pro Gly Val His Thr
130 135 140
Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser
145 150 155

<210> 7
<211> 477
<212> DNA
<213> Homosapien

<400> 7

gacatcgta tgacccagtc tccagctcc ctggctgagt ctctggcga gagggccacc 60
atcaattgca agtcagcca gagtgttta tatagctcca acaataagaa ctacttagtt 120
tggtaccagc agaaaacttagg acagccccct aagctgctca tttactggc atctaccgg 180
gaatccgggg tccctgaccg attcagtgcc agcgggtctg ggacagattt cactctcacc 240
atcagcagcc tgcaggctga agatgtggca gtttattact gtcaacaata ttatcgtagt 300
ccgtggacgt tcggccaagg gaccaaggtg gaaatcaaac gaactgtggc tgaccatct 360
gtcttcatct tcccgcacatc tcatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgcc 477

<210> 8

<211> 159

<212> PRT

<213> Homosapien

<400> 8

Asp	Ile	Val	Met	Thr	Gln	Ser	Pro	Ala	Ser	Leu	Ala	Glu	Ser	Leu	Gly
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Glu	Arg	Ala	Thr	Ile	Asn	Cys	Lys	Ser	Ser	Gln	Ser	Val	Leu	Tyr	Ser
								20				25		30	
Ser	Asn	Asn	Lys	Asn	Tyr	Leu	Val	Trp	Tyr	Gln	Gln	Lys	Leu	Gly	Gln
									40			45			
Pro	Pro	Lys	Leu	Leu	Ile	Tyr	Trp	Ala	Ser	Thr	Arg	Glu	Ser	Gly	Val
								50			55		60		
Pro	Asp	Arg	Phe	Ser	Gly	Ser	Gly	Ser	Gly	Thr	Asp	Phe	Thr	Leu	Thr
								65			70		75		80
Ile	Ser	Ser	Leu	Gln	Ala	Glu	Asp	Val	Ala	Val	Tyr	Tyr	Cys	Gln	Gln
								85			90		95		
Tyr	Tyr	Arg	Ser	Pro	Trp	Thr	Phe	Gly	Gln	Gly	Thr	Lys	Val	Glu	Ile
								100			105		110		
Lys	Arg	Thr	Val	Ala	Ala	Pro	Ser	Val	Phe	Ile	Phe	Pro	Pro	Ser	Asp
								115			120		125		
Glu	Gln	Leu	Lys	Ser	Gly	Thr	Ala	Ser	Val	Val	Cys	Leu	Leu	Asn	Asn
								130			135		140		
Phe	Tyr	Pro	Arg	Glu	Ala	Lys	Val	Gln	Trp	Lys	Val	Asp	Asn	Ala	
								145			150		155		

<210> 9

<211> 556

<212> PRT

<213> Homosapien

<400> 9

Cys	Ala	Gly	Gly	Thr	Cys	Cys	Ala	Gly	Cys	Thr	Gly	Gly	Thr	Ala	Cys
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Ala	Gly	Thr	Cys	Thr	Gly	Gly	Gly	Cys	Thr	Gly	Ala	Gly	Gly	Thr	
								20			25		30		
Gly	Ala	Ala	Gly	Ala	Ala	Gly	Cys	Cys	Thr	Gly	Gly	Gly	Cys	Cys	
								35			40		45		
Thr	Cys	Ala	Gly	Thr	Gly	Ala	Ala	Gly	Gly	Thr	Cys	Thr	Cys	Cys	Thr
								50			55		60		
Gly	Cys	Ala	Ala	Gly	Gly	Thr	Thr	Thr	Cys	Cys	Gly	Gly	Ala	Thr	Ala
								65			70		75		80
Cys	Ala	Cys	Cys	Cys	Thr	Cys	Ala	Cys	Thr	Gly	Ala	Ala	Thr	Thr	Ala

85	90	95
Thr Cys Cys Ala Thr Gly Cys Ala Cys	Thr Gly Gly Gly Thr Gly Cys	
100	105	110
Gly Ala Cys Ala Gly Gly Cys	Thr Cys Cys Thr Gly Gly Ala Ala Ala	
115	120	125
Ala Gly Gly Gly Cys Thr Thr Gly Ala Gly Thr Gly Gly Ala Thr Gly		
130	135	140
Gly Gly Ala Gly Gly Thr Thr Thr Gly Ala Thr Cys Cys Thr Gly		
145	150	155
Ala Ala Gly Ala Thr Gly Gly Thr Gly Ala Ala Ala Cys Ala Ala Thr		
165	170	175
Cys Thr Ala Cys Gly Cys Ala Cys Ala Gly Ala Ala Gly Thr Thr Cys		
180	185	190
Cys Ala Gly Gly Cys Ala Gly Ala Gly Thr Cys Ala Cys Cys Ala		
195	200	205
Thr Gly Ala Cys Cys Gly Ala Gly Gly Ala Cys Ala Cys Ala Thr Cys		
210	215	220
Thr Ala Cys Ala Gly Ala Cys Ala Cys Ala Gly Cys Cys Thr Ala Cys		
225	230	235
Ala Thr Gly Gly Ala Gly Cys Thr Gly Ala Gly Cys Ala Gly Cys Cys		
245	250	255
Thr Gly Ala Gly Ala Thr Cys Thr Gly Ala Gly Gly Ala Cys Ala Cys		
260	265	270
Gly Gly Cys Cys Gly Thr Gly Thr Ala Thr Thr Ala Cys Thr Gly Thr		
275	280	285
Gly Cys Ala Ala Cys Ala Ala Ala Cys Gly Ala Thr Thr Thr Thr Thr		
290	295	300
Gly Gly Ala Gly Thr Gly Gly Thr Ala Thr Thr Ala Thr Ala Ala		
305	310	315
Cys Thr Ala Cys Thr Gly Gly Gly Cys Cys Ala Gly Gly Ala		
325	330	335
Ala Cys Cys Cys Thr Gly Gly Thr Cys Ala Cys Cys Gly Thr Cys Thr		
340	345	350
Cys Cys Thr Cys Ala Gly Cys Cys Thr Cys Cys Ala Cys Cys Ala Ala		
355	360	365
Gly Gly Gly Cys Cys Ala Thr Cys Gly Gly Thr Cys Thr Thr Cys		
370	375	380
Cys Cys Cys Cys Thr Gly Gly Cys Cys Cys Cys Thr Gly Cys Thr		
385	390	395
Cys Cys Ala Gly Gly Ala Gly Cys Ala Cys Cys Thr Cys Cys Gly Ala		
405	410	415
Gly Ala Gly Cys Ala Cys Ala Gly Cys Gly Gly Cys Cys Cys Thr Gly		
420	425	430
Gly Gly Cys Thr Gly Cys Cys Thr Gly Gly Thr Cys Ala Ala Gly Gly		
435	440	445
Ala Cys Thr Ala Cys Thr Thr Cys Cys Cys Gly Ala Ala Cys Cys		
450	455	460
Gly Gly Thr Gly Ala Cys Gly Gly Thr Gly Thr Cys Gly Thr Gly Gly		
465	470	475
Ala Ala Cys Thr Cys Ala Gly Gly Cys Gly Cys Thr Cys Thr Gly Ala		
485	490	495
Cys Cys Ala Gly Cys Gly Gly Cys Gly Thr Gly Cys Ala Cys Ala Cys		
500	505	510
Cys Thr Thr Cys Cys Cys Ala Gly Cys Thr Gly Thr Cys Cys Thr Ala		
515	520	525

Cys	Ala	Gly	Thr	Cys	Cys	Thr	Cys	Ala	Gly	Gly	Ala	Cys	Thr	Cys	Thr
530		535						540							
Ala	Cys	Thr	Cys	Cys	Thr	Cys	Ala	Gly	Cys	Ala					
545				550				555							

<210> 10
<211> 185
<212> PRT
<213> Homosapien

<400> 10															
Gln	Val	Gln	Leu	Val	Gln	Ser	Gly	Ala	Glu	Val	Lys	Lys	Pro	Gly	Ala
1		5						10					15		
Ser	Val	Lys	Val	Ser	Cys	Lys	Val	Ser	Gly	Tyr	Thr	Leu	Thr	Glu	Leu
		20					25					30			
Ser	Met	His	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Met
	35				40						45				
Gly	Gly	Phe	Asp	Pro	Glu	Asp	Gly	Glu	Thr	Ile	Tyr	Ala	Gln	Lys	Phe
	50				55					60					
Gln	Gly	Arg	Val	Thr	Met	Thr	Glu	Asp	Thr	Ser	Thr	Asp	Thr	Ala	Tyr
65			70				75					80			
Met	Glu	Leu	Ser	Ser	Leu	Arg	Ser	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys
	85					90					95				
Ala	Thr	Asn	Asp	Phe	Trp	Ser	Gly	Tyr	Tyr	Asn	Tyr	Trp	Gly	Gln	Gly
	100					105					110				
Thr	Leu	Val	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe
	115					120					125				
Pro	Leu	Ala	Pro	Cys	Ser	Arg	Ser	Thr	Ser	Glu	Ser	Thr	Ala	Ala	Leu
	130					135					140				
Gly	Cys	Leu	Val	Lys	Asp	Tyr	Phe	Pro	Glu	Pro	Val	Thr	Val	Ser	Trp
145				150				155					160		
Asn	Ser	Gly	Ala	Leu	Thr	Ser	Gly	Val	His	Thr	Phe	Pro	Ala	Val	Leu
					165			170					175		
Gln	Ser	Ser	Gly	Leu	Tyr	Ser	Leu	Ser							
				180			185								

<210> 11
<211> 490
<212> DNA
<213> Homosapien

<400> 11															
gacatcgta	tgacccagtc	tccagactcc	ctggctgtgt	ctctgggcga	gagggccacc	60									
atcaactgca	agtccagcca	gagtgtttta	tacagctcca	acaataagaa	ctacttagtt	120									
tggtaccaac	agaaaccagg	acagcctcct	aaactgctca	tttactgggc	atctatccgg	180									
gaatccgggg	tccctgaccg	attcagtgcc	agcgggtctg	ggacagattt	cactctcacc	240									
atcaacagcc	tgcaggctga	agatgtggca	gttattact	gtcagcagta	tttttatagt	300									
ccgtggacgt	tcggccaagg	gaccaaggtg	gaaatcaaac	gaactgtggc	tgcaccatct	360									
gtcttcatct	tcccgccatc	tcatgagcag	ttgaaatctg	gaactgcctc	tgttgtgc	420									
ctgctgaata	acttctatcc	cagagaggcc	aaagtacagt	ggaaggtgga	taacgcccctc	480									
caatcggtta						490									

<210> 12

<211> 163
<212> PRT
<213> Homosapien

<400> 12
Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
1 5 10 15
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
20 25 30
Ser Asn Asn Lys Asn Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln
35 40 45
Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Ile Arg Glu Ser Gly Val
50 55 60
Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65 70 75 80
Ile Asn Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
85 90 95
Tyr Phe Tyr Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
100 105 110
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
115 120 125
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
130 135 140
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu
145 150 155 160
Gln Ser Gly

<210> 13
<211> 543
<212> DNA
<213> Homosapien

<400> 13
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tcctgcagg tttccggaca caccctact gaattatcca tgcactgggt ggcacaggct 120
cctggaaaag ggcttgagtg gatgggaggt tttgatcctg aagatgtga aacaatctac 180
gcacagaagt tccaggacag agtcaccatg accgaggaca catctacaga cacagcctac 240
atggagctga gcagccctaaatctgaggac acggccgtgt attactgtgc aaccaacgtat 300
ttttggagtg gttatatttga ctgctggggc cagggaaaccc tggtcaccgt ctccctagcc 360
tccaccaagg gccccatcggt ctccccctg gcccctgtt ccaggagcac ctcccgagagc 420
acagcggccc tgggctgcct ggtcaaggac tactccccg aaccgggtgac ggtgtcgtgg 480
aactcaggcg ctctgaccag cggcgtgcac accttcccgat ctgtcctaca gtcctcagga 540
ctt 543

<210> 14
<211> 181
<212> PRT
<213> Homosapien

<400> 14
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly His Thr Leu Thr Glu Leu

20	25	30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met		
35	40	45
Gly Gly Phe Asp Pro Glu Asp Asp Glu Thr Ile Tyr Ala Gln Lys Phe		
50	55	60
Gln Asp Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr		
65	70	75
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys		
85	90	95
Ala Thr Asn Asp Phe Trp Ser Gly Tyr Phe Asp Cys Trp Gly Gln Gly		
100	105	110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe		
115	120	125
Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu		
130	135	140
Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp		
145	150	155
Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu		
165	170	175
Gln Ser Ser Gly Leu		
180		

<210> 15
 <211> 490
 <212> DNA
 <213> Homosapien

<400> 15
 gacatcggtgc tgaccaggc tccagactcc ctggctgtgt gtctggcgaa gaggccacc 60
 atcaactgca agtccaggcc aagtgtttta tatagtccca acaataagaa cttcttagtt 120
 tggtaccaggc agagaccagg acagccctt aagctgctca tttactgggc atctaccgg 180
 gaatccgggg tccctgaccg attcagtgcc agcgggtctg ggacagattt cactctcacc 240
 atcagcagcc tgcaggctga agatgtggca gtttattact gtcagcaata ttatagtagt 300
 ccgtggacgt tcggccaagg gaccaagggtg gaaatcaaac gaaactgtggc tgaccatct 360
 gtcttcatct tcccgccatc tgatgagcag ttgaaatctg gaaactgcctc tgggtgtgc 420
 ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgccctc 480
 caatcggtt 490

<210> 16
 <211> 163
 <212> PRT
 <213> Homosapien

<400> 16
 Asp Ile Val Leu Thr Gln Ser Pro Asp Ser Leu Ala Val Cys Leu Gly
 1 5 10 15
 Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
 20 25 30
 Pro Asn Asn Lys Asn Phe Leu Val Trp Tyr Gln Gln Arg Pro Gly Gln
 35 40 45
 Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
 50 55 60
 Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
 65 70 75 80

<210> 17
<211> 1335
<212> DNA
<213> Homosapien

<400> 17
caggtccagc tggtacagtc tggggcttag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcaagg ttccggata caccctcact gaattatcca tgcactgggt gcgacagggc 120
cctggaaaag ggcttgagtg gatgggaggt tttgatccctg aagatggtga aacaatctac 180
gcacagaagt tccagggcag agtcaccatg accgaggaca catctacaga cacagtctac 240
atggagctga gcagcctgag atctgaggac acggccatgt attactgtgc aacacggag 300
tttggactg gtatTTGA ccactggggc cagggAACCC tggtaccgt ctcctcagcc 360
tccaccaagg gcccattcggt cttcccccgt ggcgcctgtct ccaggagcac ctccgagagc 420
acagcggccc tgggctgcct ggtcaaggac tacttccccg aaccgggtgac ggtgtcggtgg 480
aactcaggcg ctctgaccag cggcgtgcac accttcccag ctgtcctaca gtcctcagga 540
ctctactccc tcagcagcgt ggtgaccgtg ccctccagca acttcggcac ccagacccatc 600
acctgcaacg tagatcacaa gcccagcaac accaagggtgg acaagacagt tgagcgcaaa 660
tgggtgtcg agtgcaccacc gtgcccagca ccacctgtgg caggaccgtc agtcttcctc 720
ttccccccaa aacccaaggaa caccctcatg atctcccgaa cccctgaggt cacgtcgctg 780
gtggtggacg tgagccacga agaccccggag gtccaggatca actgttacgt ggacggcgtg 840
gaggtgcata atgccaagac aaagccacgg gaggagcaatgt tcaacagcac gttccgtgtg 900
gtcagcgtcc tcaccgttgt gcaccaggac tggctgaacg gcaaggagta caagtgcag 960
gtctccaaaca aaggcctccc agccccatc gagaaaaacca tctccaaaac caaagggcag 1020
ccccgagaac cacaggtgtc caccctgccc ccatccccggg aggagatgac caagaaccag 1080
gtcagcctga cctgccttgtt caaaggcttc taccggcagcg acatcgccgt ggagtgggag 1140
agcaatgggc agccggagaa caactacaag accacacaccc ccatgtcgga ctccgacggc 1200
tccttcttcc tctacagcaa gctcaccgtg gacaagagca ggtggcagca gggggAACGTC 1260
ttctcatgtt ccgtgtatgc tggggctctg cacaaccact acacgcagaa gagcctctcc 1320
ctgtctccgg gtaaa 1335

<210> 18
<211> 445
<212> PRT
<213> Homosapien

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<400> 18
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
  1           5           10           15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
  20          25          30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met

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35	40	45													
Gly	Gly	Phe	Asp	Pro	Glu	Asp	Gly	Glu	Thr	Ile	Tyr	Ala	Gln	Lys	Phe
50					55					60					
Gln	Gly	Arg	Val	Thr	Met	Thr	Glu	Asp	Thr	Ser	Thr	Asp	Thr	Val	Tyr
65					70					75					80
Met	Glu	Leu	Ser	Ser	Leu	Arg	Ser	Glu	Asp	Thr	Ala	Met	Tyr	Tyr	Cys
					85					90					95
Ala	Thr	Arg	Glu	Phe	Trp	Thr	Gly	Tyr	Phe	Asp	His	Trp	Gly	Gln	Gly
					100					105					110
Thr	Leu	Val	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe
					115					120					125
Pro	Leu	Ala	Pro	Cys	Ser	Arg	Ser	Thr	Ser	Glu	Ser	Thr	Ala	Ala	Leu
					130					135					140
Gly	Cys	Leu	Val	Lys	Asp	Tyr	Phe	Pro	Glu	Pro	Val	Thr	Val	Ser	Trp
145					150					155					160
Asn	Ser	Gly	Ala	Leu	Thr	Ser	Gly	Val	His	Thr	Phe	Pro	Ala	Val	Leu
					165					170					175
Gln	Ser	Ser	Gly	Leu	Tyr	Ser	Leu	Ser	Ser	Val	Val	Thr	Val	Pro	Ser
					180					185					190
Ser	Asn	Phe	Gly	Thr	Gln	Thr	Tyr	Thr	Cys	Asn	Val	Asp	His	Lys	Pro
					195					200					205
Ser	Asn	Thr	Lys	Val	Asp	Lys	Thr	Val	Glu	Arg	Lys	Cys	Cys	Val	Glu
					210					215					220
Cys	Pro	Pro	Cys	Pro	Ala	Pro	Pro	Val	Ala	Gly	Pro	Ser	Val	Phe	Leu
225					230					235					240
Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg	Thr	Pro	Glu
					245					250					255
Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro	Glu	Väl	Gln
					260					265					270
Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala	Lys	Thr	Lys
					275					280					285
Pro	Arg	Glu	Glu	Gln	Phe	Asn	Ser	Thr	Phe	Arg	Val	Val	Ser	Val	Leu
					290					295					300
Thr	Val	Val	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr	Lys	Cys	Lys
305					310					315					320
Val	Ser	Asn	Lys	Gly	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr	Ile	Ser	Lys
					325					330					335
Thr	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu	Pro	Pro	Ser
					340					345					350
Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys	Leu	Val	Lys
					355					360					365
Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser	Asn	Gly	Gln
					370					375					380
Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Met	Leu	Asp	Ser	Asp	Gly
385					390					395					400
Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser	Arg	Trp	Gln
					405					410					415
Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala	Leu	His	Asn
					420					425					430
His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys			
					435					440					445

<210> 19
<211> 660

<212> DNA
<213> Homosapien

<400> 19
gacatcgtga tgaccaggc tccagactcc ctggctgtgt ctctggcga gagggccacc 60
atcaactgca agtccagcca gagtgttta tacagctcca acaataagaa ctacttagtt 120
tggtatcagc agaaaaccagg acagcctctt aaactgctca ttactggc atctatccgg 180
gaatccgggg tcccgaccc attcagtgcc agccggctcg ggacagattt cactctcacc 240
atcagcagcc tgcaggctga agatgtgca gtttattact gtcagcaata ttatagtact 300
ccgctcactt tcggcggagg gaccaagggtg gagatcaaac gaactgtggc tgacccatct 360
gtcttcatct tcccgccatc tgatgagcag ttgaaatctg gaactgcctc tggtgtgtgc 420
ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaaggtgga taacgcctc 480
caatcgggta actcccaagga gagtgtcaca gagcaggaca gcaaggacag cacctacagc 540
ctcagcagca ccctgacgct gagcaaagca gactacgaga aacacaaagt ctacgcctgc 600
gaagtcaccc atcagggcct gagctcgccc gtcacaaaga gcttcaacag gggagagtgt 660

<210> 20
<211> 220
<212> PRT
<213> Homosapien

<400> 20
Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
1 5 10 15
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
20 25 30
Ser Asn Asn Lys Asn Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln
35 40 45
Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Ile Arg Glu Ser Gly Val
50 55 60
Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65 70 75 80
Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
85 90 95
Tyr Tyr Ser Thr Pro Leu Thr Phe Gly Gly Thr Lys Val Glu Ile
100 105 110
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
115 120 125
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
130 135 140
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu
145 150 155 160
Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp
165 170 175
Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr
180 185 190
Glu Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser
195 200 205
Ser Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
210 215 220

<210> 21
<211> 543

<212> DNA
<213> Homosapien

<400> 21
caggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcaagg tttccggata cactttact gaattatcca tgcactgggt gcgacaggct 120
cctggaaaag ggcttgagtg gatgggaggt tttgatcctg aagatggtga aacaagctac 180
gcacagaagt tccggggcag agtcaccatg accgaggaca catctacaga cacagccac 240
atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aaccaacgat 300
ttttggagtg gttatgttga ctattgggc cagggAACCC tggtcaccgt ctcctcagcc 360
tccaccaagg gcccattcggt ctccccctg gcgcctgct ccaggagcac ctccgagagc 420
acagcggccc tgggctgcct ggtcaaggac tactccccg aaccggtgac ggtgtcgtgg 480
aactcaggcg ctctgaccag cggcgtgcac accttccagc ctgtcctaca gtcctcagga 540
ctt 543

<210> 22
<211> 181
<212> PRT
<213> Homosapien

<400> 22
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Phe Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ser Tyr Ala Gln Lys Phe
50 55 60
Arg Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala His
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr Asn Asp Phe Trp Ser Gly Tyr Phe Asp Tyr Trp Gly Gln Gly
100 105 110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125
Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
130 135 140
Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
145 150 155 160
Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu
165 170 175
Gln Ser Ser Gly Leu
180

<210> 23
<211> 460
<212> DNA
<213> Homosapien

<400> 23
gacatccaga tgaccaggatc tccatcttcc gtgtctgcat ctgttaggaga cagagtcacc 60
atcacttgc gggcgagtca gggattgac atctacttag cctggatca gcagaaacca 120

gggaaagccc ctaagctcct gatcaatgct gcatccagg ttgcaaaacgg ggtccccctca 180
aggttcggcg gcagtggatc tgggacagat ttcaactctca ccatcagcgg cctgcagcct 240
gaagattttg caacttacta ttgtcaactg acttactttt tcccgtggac gttcggccaa 300
gggaccaagg tggaaatcaa acgaactgtg gctgcaccat ctgtcttcat ctccccgcca 360
tctgatgagc agttgaaatc tggaaactgccc tctgttgtgt gcctgctgaa taacttctat 420
cccagagagg ccaaagtaca gtggaaagggtg gataacgccc 460

<210> 24
<211> 153
<212> PRT
<213> Homosapien

<400> 24
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Val Ser Ala Ser Val Gly
1 5 10 15
Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Asp Ile Tyr
20 25 30
Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
35 40 45
Asn Ala Ala Ser Ser Leu Gln Asn Gly Val Pro Ser Arg Phe Gly Gly
50 55 60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Gly Leu Gln Pro
65 70 75 80
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Leu Thr Tyr Phe Phe Pro Trp
85 90 95
Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala Ala
100 105 110
Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly
115 120 125
Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala
130 135 140
Lys Val Gln Trp Lys Val Asp Asn Ala
145 150

<210> 25
<211> 543
<212> DNA
<213> Homosapien

<400> 25
caggtccagc tggcacagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcaagg ttccggata caccctact gaattatcca tgcactgggt gcgacgaatt 120
cctggaaaag ggcttgagtg gatgggaggt tttgaccctg aagatggta aacaatctac 180
gcacagaagt tccagggcag agtcaccatg accgaggaca catctacaga cacagcctac 240
atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aacaaacgat 300
ttttggagtg gctattgggg ccactggggc cagggAACCC tggtcaccgt ctccctcagcc 360
tccaccaagg gcccattcggt cttcccccgt gcgcctgtc ccaggagcac ctccgagagc 420
acagcggccc tgggctgcct ggtcaaggac tacttccccg aaccgggtgac ggtgtcgtgg 480
aactcaggcgc ctctgaccag cggcgtgcac accttcccag ctgtcctaca gtcctcagga 540
ctt 543

<210> 26
<211> 181
<212> PRT

<213> Homosapien

<400> 26

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Arg Ile Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr Asn Asp Phe Trp Ser Gly Tyr Trp Gly His Trp Gly Gln Gly
100 105 110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125
Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
130 135 140
Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
145 150 155 160
Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu
165 170 175
Gln Ser Ser Gly Leu
180

<210> 27

<211> 459

<212> DNA

<213> Homosapien

<400> 27

gacatcgtga tgaccaggc tccagactcc ctggctgtgt ctctggcgaa gaggggccacc 60
atcaactgca agtccagccaa gagtgttta tacagctcca acaataagaa ctaccttagct 120
tggcaccaag ctgctcattt actggacata tatccggaa tccgggtcc ctgaccgatt 180
cagttggcaggc gggctctggaa cagatttcac tctcaccatc agcagcctgc aggctgaaga 240
tgtggcagtt tattactgtc aggaacattha tagtattccg tggacgttgc gccaaggggac 300
caaggtggaa atcaaacgaa ctgtggctgc accatctgtc ttcatcttcc cgccatctga 360
tgagcagttt aactgcctct gttgtgtgcc tgctgaataa cttctatccc agagaggcca 420
aagtacagtgaaggtggat aacggccctcc aatcgggtaa 459

<210> 28

<211> 149

<212> PRT

<213> Homosapien

<400> 28

Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
1 5 10 15
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
20 25 30
Ser Asn Asn Lys Asn Tyr Leu Ala Trp Tyr Leu Leu Ile Tyr Trp Thr

35	40	45	
Tyr Ile Arg Glu Ser Gly Val Pro Asp Arg Phe Ser Gly Ser			
50	55	60	
Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Ala Glu Asp Val			
65	70	75	80
Ala Val Tyr Tyr Cys Gln Glu His Tyr Ser Ile Pro Trp Thr Phe Gly			
85	90	95	
Gln Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala Ala Pro Ser Val			
100	105	110	
Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Asn Cys Leu Cys Cys Val			
115	120	125	
Pro Ala Glu Leu Leu Ser Gln Arg Gly Gln Ser Thr Val Glu Gly Gly			
130	135	140	
Arg Pro Pro Ile Gly			
145			

<210> 29
<211> 524
<212> DNA
<213> Homosapien

<400> 29
caggtccagc tggcacagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcagg tttccggata caccctact gaattatcca tgcactgggt gcgacaggct 120
cctggaaaag ggcttgagtg gatgggaggt tttgatcctg aagatgatga aacaatctac 180
gcacagaagt tccagggcag agtcaccatg accgaggaca catctacaga cacggcctac 240
atggagctga gcagcctgag atctgaggac acggccgtgt atttctgtgc aaccaacgat 300
ttttggagtg gttatattga ctgctggac cagggAACCC tggtcaccgt ctccctcagcc 360
tccaccaagg gcccacatcggt ctccccctg gcgcctgtct ccaggaacac ctccgagagc 420
acagcggccc tgggctgcct ggtcaaggac tacttccccg aaccggtgac ggtgtcgtgg 480
aactcaggcg ctctgaccag cggcgtgcac accttcccag ctgt 524

<210> 30
<211> 174
<212> PRT
<213> Homosapien

<400> 30
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Asp Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Phe Cys
85 90 95
Ala Thr Asn Asp Phe Trp Ser Gly Tyr Phe Asp Cys Trp Asp Gln Gly
100 105 110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125

Pro Leu Ala Pro Cys Ser Arg Asn Thr Ser Glu Ser Thr Ala Ala Leu
 130 135 140
 Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
 145 150 155 160
 Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala
 165 170

<210> 31
 <211> 490
 <212> DNA
 <213> Homosapien

<400> 31
 gacatcgta tgacccagtc tccagactcc ctggctgcgt ctctggcga gagggccacc 60
 atcaactgca agtccagtca gagtgtttt acaataagaa ttatggtagtt 120
 tggtaccagg aaaaaccagg acagcctcct aagctgctca tttactggc atctatccgg 180
 gaatccgggg tccctgaccg attcagtgcc agcgggtctg ggacagattt cactctcacc 240
 atcagcagcc tgcaggctga agatgtggca gtttatttct gtcagcaata ttatagttct 300
 ccgtggacgt ttggccaagg gaccaaggtg gaaatcaaac gaactgtggc tgcaccatct 360
 gtcttcatct tcccgccatc tgatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
 ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgcctc 480
 caatcggtta 490

<210> 32
 <211> 163
 <212> PRT
 <213> Homosapien

<400> 32
 Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Ala Ser Leu Gly
 1 5 10 15
 Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Arg
 20 25 30
 Ser Asn Asn Lys Asn Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln
 35 40 45
 Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Ile Arg Glu Ser Gly Val
 50 55 60
 Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
 65 70 75 80
 Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Phe Cys Gln Gln
 85 90 95
 Tyr Tyr Ser Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
 100 105 110
 Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
 115 120 125
 Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
 130 135 140
 Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu
 145 150 155 160
 Gln Ser Gly

<210> 33

<211> 545
<212> DNA
<213> Homosapien

<400> 33
caggtccagc tggcacagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggtc 60
tcctgcaagg tttccggata caccctcaact gaattatcca tgcactgggt gcgacaggct 120
cctggaaaag ggcttgagtg gatgggaggt tttgatcctg aagatggtga aacaatctac 180
gcacagaagt tccagggcag agtcaccatg accgaggaca catctacaga cacagcctac 240
atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aacctggtat 300
agtggatct acttagctt tgatatctgg ggc当地ggca 360
gcctccacca agggccatc ggtcttcccc ctggcgcct gctccaggag cacctccgag 420
agcacagcgg ccctgggctg cctggtaag gactacttcc ccgaaccggt gacgggtcg 480
tggactcag ggc当地ctgac cagcggcgtg cacaccttcc cagctgtcct acagtccctca 540
ggatt 545

<210> 34
<211> 181
<212> PRT
<213> Homosapien

<400> 34
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr Trp Tyr Ser Gly Ile Tyr Leu Ala Phe Asp Ile Trp Gly Gln
100 105 110
Gly Thr Met Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val
115 120 125
Phe Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala
130 135 140
Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser
145 150 155 160
Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val
165 170 175
Leu Gln Ser Ser Gly
180

<210> 35
<211> 472
<212> DNA
<213> Homosapien

<400> 35
gaaattgtgc tgactcagtc tccagacttt cagtcgtga ctccaaagga gaaagtccacc 60

atcacctgcc gggccagtca gagcatttgtt agtagcttac actggatcca gcagaaacca 120
 gatcagtctc caaagctctt catcaagttt gcttcccagt cttctcagg ggtcccctcg 180
 aggttcagtgc cagtggtatc tgggacatgtt cttaccctca ccatcaatag cctggaaagct 240
 gaagatgctg caacgttata ctgtcatcg agtagtagtt tacctcacac tttcggcgga 300
 gggaccaagg tggagatcaa acgaactgtg gctgcaccat ctgtcttcat cttcccgcca 360
 tctgatgagc agttgaaatc tggaaactgtc tcttgtgtgt gcctgctgaa taacttctat 420
 cccagagagg ccaaagtaca gtggaaagggtg gataacgccc tccaaatcgaa ta 472

<210> 36
 <211> 157
 <212> PRT
 <213> Homosapien

<400> 36
 Glu Ile Val Leu Thr Gln Ser Pro Asp Phe Gln Ser Val Thr Pro Lys
 1 5 10 15
 Glu Lys Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Gly Ser Ser
 20 25 30
 Leu His Trp Tyr Gln Gln Lys Pro Asp Gln Ser Pro Lys Leu Leu Ile
 35 40 45
 Lys Tyr Ala Ser Gln Ser Phe Ser Gly Val Pro Ser Arg Phe Ser Gly
 50 55 60
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Asn Ser Leu Glu Ala
 65 70 75 80
 Glu Asp Ala Ala Thr Tyr Tyr Cys His Gln Ser Ser Ser Leu Pro His
 85 90 95
 Thr Phe Gly Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala Ala
 100 105 110
 Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly
 115 120 125
 Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala
 130 135 140
 Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly
 145 150 155

<210> 37
 <211> 1335
 <212> DNA
 <213> Homosapien

<400> 37
 caggtccagt tggcacatc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
 tcctgcaagg tttccggata caccctactt gaattatcca tgcactgggt gcgacaggct 120
 cctggaaaag ggcttgagtgc gatgggaggt tttgatcctg aagatggtga aacaatctac 180
 gcacagaagt tccagggcag agtcagtatg accgaggaca catccacaga cacagcctac 240
 atggagctga gcagcctgatg atctgaggac acggccgtgt atttctgtgc aaccaacgaa 300
 ttttggagtgc gttatatttgc ctactggggc cagggaaaccc tggtcaccgt ctccctcagcc 360
 tccaccaagg gcccattcggt ctccccctg gcccctgtt ccaggagcac ctcccgagac 420
 acagcggccc tgggctgcct ggtcaaggac tacttccccg aaccggtgac ggtgtcggtgg 480
 aactcaggcg ctctgaccat cggcgtgcac accttcccag ctgtcctaca gtcctcagga 540
 ctctactccc tcagcagcgt ggtgaccgtg ccctccagca acttcggcac ccagacctac 600
 acctgcaacg tagatcacaa gcccagcaac accaagggtgg acaagacagt tgagcgaaa 660
 tgggtgtcg agtggcccacc gtggcccgatca ccacctgtgg caggaccgtc agtcttcctc 720
 ttccccccaa aacccaagga caccctcatg atctcccgaa cccctgaggt cacgtgcgtg 780

gtggggacg tgagccacga agaccccgag gtccagttca actggtagt ggacggcgtg 840
 gaggtgcata atgccaagac aaagccacgg gagggcagt tcaacagcac gttccgtgtg 900
 gtcagcgtcc tcacccgtgt gcaccaggac tggctgaacg gcaaggagta caagtgcac 960
 gtctccaaca aaggccccc agccccatc gagaaaaacca tctccaaaac caaaggccag 1020
 ccccgagaac cacaggtgt a caccctgccc ccatcccgagg aggagatgac caagaaccag 1080
 gtcagcctga cctgcctgtt caaaggcttc taccggcagcg acatggccgt ggagtggag 1140
 agcaatgggc agccggagaa caactacaag accacaccc ccatgctgga ctccgacggc 1200
 tccttcttcc tctacagcaa gtcaccgtg gacaagagca ggtggcagca gggaaacgtc 1260
 ttctcatgtt ccgtgatgca tgaggctctg cacaaccact acacgcagaa gagcctctcc 1320
 ctgtctccgg gtaaa 1335

<210> 38
 <211> 445
 <212> PRT
 <213> Homosapien

<400> 38
 Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
 1 5 10 15
 Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
 20 25 30
 Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
 35 40 45
 Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
 50 55 60
 Gln Gly Arg Val Ser Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
 65 70 75 80
 Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Phe Cys
 85 90 95
 Ala Thr Asn Glu Phe Trp Ser Gly Tyr Phe Asp Tyr Trp Gly Gln Gly
 100 105 110
 Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
 115 120 125
 Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
 130 135 140
 Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
 145 150 155 160
 Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu
 165 170 175
 Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser
 180 185 190
 Ser Asn Phe Gly Thr Gln Thr Tyr Thr Cys Asn Val Asp His Lys Pro
 195 200 205
 Ser Asn Thr Lys Val Asp Lys Thr Val Glu Arg Lys Cys Cys Val Glu
 210 215 220
 Cys Pro Pro Cys Pro Ala Pro Pro Val Ala Gly Pro Ser Val Phe Leu
 225 230 235 240
 Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu
 245 250 255
 Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Gln
 260 265 270
 Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys
 275 280 285
 Pro Arg Glu Glu Gln Phe Asn Ser Thr Phe Arg Val Val Ser Val Leu
 290 295 300

Thr Val Val His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys
 305 310 315 320
 Val Ser Asn Lys Gly Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys
 325 330 335
 Thr Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser
 340 345 350
 Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys
 355 360 365
 Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln
 370 375 380
 Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Met Leu Asp Ser Asp Gly
 385 390 395 400
 Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln
 405 410 415
 Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn
 420 425 430
 His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
 435 440 445

<210> 39
 <211> 660
 <212> DNA
 <213> Homosapien

<400> 39
 gacatcgta tgacccagtc tccagactcc ctggctgtgt ctctggcga gagggccacc 60
 atcaactgca agtccagcca gagtgttta tacagctcca acaataagaa ctattnatgtt 120
 tggtaccagc agagaccagg acagcctctt aagctgctca ttactggc atctaccgg 180
 gaatccgggg tccctgaccg attcagtgcc agccggcttg ggacagattt cactctcacc 240
 atcagcagcc tgcaggctga agatgtggca gtttattact gtcagcaata ttttattct 300
 ccgtggacgt tcggccaagg gaccaaggta gaaatcaaac gaaactgtggc tgcaccatct 360
 gtcttcatct tccccccatc tcatgagcag ttgaaatctg gaaactgcctc tgggtgtgc 420
 ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgcctc 480
 caatcgggta actcccaagga gagtgcaca gagcaggaca gcaaggacag cacctacago 540
 ctcagcagca ccctgacgct gagcaaagca gactacgaga aacacaagt ctacgcctgc 600
 gaagtcaccc atcagggcct gagctcgccc gtcacaaaga gcttcaacag gggagagtgt 660

<210> 40
 <211> 220
 <212> PRT
 <213> Homosapien

<400> 40
 Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
 1 5 10 15
 Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
 20 25 30
 Ser Asn Asn Lys Asn Tyr Leu Val Trp Tyr Gln Gln Arg Pro Gly Gln
 35 40 45
 Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
 50 55 60
 Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
 65 70 75 80

Ile	Ser	Ser	Leu	Gln	Ala	Glu	Asp	Val	Ala	Val	Tyr	Tyr	Cys	Gln	Gln
85									90					95	
Tyr	Phe	Tyr	Ser	Pro	Trp	Thr	Phe	Gly	Gln	Gly	Thr	Lys	Val	Glu	Ile
	100						105					110			
Lys	Arg	Thr	Val	Ala	Ala	Pro	Ser	Val	Phe	Ile	Phe	Pro	Pro	Ser	Asp
115							120					125			
Glu	Gln	Leu	Lys	Ser	Gly	Thr	Ala	Ser	Val	Val	Cys	Leu	Leu	Asn	Asn
130						135					140				
Phe	Tyr	Pro	Arg	Glu	Ala	Lys	Val	Gln	Trp	Lys	Val	Asp	Asn	Ala	Leu
145						150				155			160		
Gln	Ser	Gly	Asn	Ser	Gln	Glu	Ser	Val	Thr	Glu	Gln	Asp	Ser	Lys	Asp
	165						170					175			
Ser	Thr	Tyr	Ser	Leu	Ser	Ser	Thr	Leu	Thr	Leu	Ser	Lys	Ala	Asp	Tyr
	180						185					190			
Glu	Lys	His	Lys	Val	Tyr	Ala	Cys	Glu	Val	Thr	His	Gln	Gly	Leu	Ser
	195						200				205				
Ser	Pro	Val	Thr	Lys	Ser	Phe	Asn	Arg	Gly	Glu	Cys				
	210					215				220					

<210> 41
<211> 556
<212> DNA
<213> Homosapien

<400> 41
caggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggtc 60
tcctgcagg tttccggaca cattttcaact gaatttatcca tacactgggt gcgacaggct 120
cctggaaaag ggctcgagtg gatggggaggt tttatccctg aagatggtga aacaatctac 180
gcacagaagt tccaggcag agtcaccatg accaggagaca catctacaga cacagtctac 240
atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aaccaacgat 300
ttttggagtg gttatatttga ctactggggc cagggaaaccc tggtcaccgt ctccctcagcc 360
tccaccaagg gcccattcggt cttcccccgt gcgccctgct ccaggagcac ctccgagagc 420
acagcggccc tgggctgcct ggtcaaggac tacttccccg aaccggtgac ggtgtcgtgg 480
aactcaggcg ctctgaccag cggcgtgcac accttcccag ctgtcctaca gtcctcagga 540
ctctactccc tcagca 556

<210> 42
<211> 185
<212> PRT
<213> Homosapien

<400> 42
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly His Ile Phe Thr Glu Leu
20 25 30
Ser Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Val Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Thr Asn Asp Phe Trp Ser Gly Tyr Phe Asp Tyr Trp Gly Gln Gly
 100 105 110
 Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
 115 120 125
 Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
 130 135 140
 Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
 145 150 155 160
 Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu
 165 170 175
 Gln Ser Ser Gly Leu Tyr Ser Leu Ser
 180 185

<210> 43
 <211> 490
 <212> DNA
 <213> Homosapien

<400> 43
 gacatcgta tgacccagtc tccaggctcc ctggctgtgt ctctggcga gagggccacc 60
 atcaactgca agtccagcca gagtattta ttcaaggcca acaataagaa ctatctaact 120
 tggtaccagc agaaaaccagg acagcctcct aaactgctca ttactggc atctatccgg 180
 gaatccgggg tccctgatcg attcagtgcc agccggctcg ggtcaaattt cactctcacc 240
 atcaccagcc tgcaggctga agatgtggca atttattact gtcagcaata ttatagtagt 300
 ccgtggacgt tcggccaagg gaccaagggtg gaaatcaaac gaactgtggc tgaccatct 360
 gtcttcatct tccccccatc tgatgagcag ttgaaatctg gaactgcctc tggtgtgtgc 420
 ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgccctc 480
 caatcgggta 490

<210> 44
 <211> 163
 <212> PRT
 <213> Homosapien

<400> 44
 Asp Ile Val Met Thr Gln Ser Pro Gly Ser Leu Ala Val Ser Leu Gly
 1 5 10 15
 Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Ile Leu Phe Arg
 20 25 30
 Ser Asn Asn Lys Asn Tyr Leu Thr Trp Tyr Gln Gln Lys Pro Gly Gln
 35 40 45
 Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Ile Arg Glu Ser Gly Val
 50 55 60
 Pro Asp Arg Phe Ser Gly Ser Gly Ser Asn Phe Thr Leu Thr
 65 70 75 80
 Ile Thr Ser Leu Gln Ala Glu Asp Val Ala Ile Tyr Tyr Cys Gln Gln
 85 90 95
 Tyr Tyr Ser Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
 100 105 110
 Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
 115 120 125
 Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
 130 135 140
 Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu

145
Gln Ser Gly

150

155

160

<210> 45
<211> 559
<212> DNA
<213> Homosapien

<400> 45
caggtccagc tggcacagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggtc 60
tcctgcaagg tttccggata caccctcaact gaattatcca tgcactgggt gcgacaggct 120
cctggaaaag ggcttgagtg gatgggaggt tttgatcctg aagatggtga aacaatcaac 180
gcacagaagt tccaggcag agtcaccatg accgaggaca catctacaga cacaggctac 240
atggagctga gcagccctgat atctgaggac acggccgtgt attactgtgc aacagatcct 300
ggtggatata gtggctactt tgaccactgg ggcaggaa ccctggtcac cgtctcctca 360
gcctccacca agggcccatc ggtttcccc ctggcgcctt gctccaggag cacctccgag 420
agcacagcgg ccctgggctg cctggtcaag gactacttcc ccgaaccggg gacgggtgtcg 480
tggactctag ggcgtctgac cagcggcgtg cacacccctcc cagctgtcct acagtccctca 540
ggactctact ccctcagca 559

<210> 46
<211> 186
<212> PRT
<213> Homosapien

<400> 46
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Asn Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Gly Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr Asp Pro Gly Gly Tyr Ser Gly Tyr Phe Asp His Trp Gly Gln
100 105 110
Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val
115 120 125
Phe Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala
130 135 140
Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser
145 150 155 160
Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val
165 170 175
Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser
180 185

<210> 47

<211> 464
<212> DNA
<213> Homosapien

<400> 47
gacatcgta tgacccagtc tccagattc ctggctgtgt ctctggcga gaggcccacc 60
atcaactgca agtccagcca gagtgtttt tacagctcca acaataagaa ctacttagtt 120
tggtaccagc agaaacccgg acagcctcct aagctgctcc tttactggc atctacccgg 180
gaatccgggg tccctgaccg attcagtgcc agcgggtctg ggacagattt cactctcacc 240
atcagcagcc tgcaggctga agatgtgca gtttattact gtcagcaata ttatagttct 300
ccgtggacgt tcggccaagg gaccaaggtg gaaatcaaac gaactgtggc tgaccatct 360
gtcttcatct tcccgcattc tgatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
ctgctgaata acttctatcc cagagagcc aaagtacagt ggaa 464

<210> 48
<211> 154
<212> PRT
<213> Homosapien

<400> 48
Asp Ile Val Met Thr Gln Ser Pro Asp Phe Leu Ala Val Ser Leu Gly
1 5 10 15
Glu Arg Pro Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Phe Tyr Ser
20 25 30
Ser Asn Asn Lys Asn Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln
35 40 45
Pro Pro Lys Leu Leu Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
50 55 60
Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65 70 75 80
Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
85 90 95
Tyr Tyr Ser Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
100 105 110
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
115 120 125
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
130 135 140
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp
145 150

<210> 49
<211> 476
<212> DNA
<213> Homosapien

<400> 49
caggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcaagg ttccggata caccctact gaattatcca tgcactgggt gcgcacaggct 120
cctggaaaag ggcttgatgt gatggaggt tttatccctg aagatgtga aacaatctac 180
gcacagaagt tccagggcag agtcaccatg accgaggaca catctacaca cacagcctac 240
atggaaactga gcagcctgag atctgaggac acggccgtgt attactgtgc aacacacgt 300
ttttggatgt cttatccatc ctactgggc cagggaaacc tggtcaccgt ctcctcagct 360
tccaccaagg gcccattccgt cttcccccgt gcgcctgtgt ccaggagcac ctccgagagc 420

acagccgccc tgggctgcct ggtcaaggac tacttccccg aaccggtgac ggtgtc 476

<210> 50
<211> 158
<212> PRT
<213> Homosapien

<400> 50
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Asp Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr His Thr Ala Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr His Asp Phe Trp Ser Ala Tyr Phe Tyr Tyr Trp Gly Gln Gly
100 105 110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125
Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
130 135 140
Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val
145 150 155

<210> 51
<211> 490
<212> DNA
<213> Homosapien

<400> 51
gacatcgta tgacccagtc tccagactcc ctggctgtgt ctctggcga gagggccacc 60
atcaactgca agtccagcca gagtgttta tacggctcca acaataagag ctacttagct 120
tggtaccaggc agaaaaccagg acagccctct aagctgctca tttactggc atctaccgg 180
gaatccgggg tccctgaccg attcagtgcc agcgggtctg ggacagattt cactctcacc 240
atcagcagcc tgcaggctgc agatgtgca gtttattact gtcagcaaca ttatagact 300
ccgtgcagtt ttggccaggg gaccaaactg gagatcaaac gaactgtggc tgaccatct 360
gtcttcatct tccccccatc tgatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgcccctc 480
caatcggtta 490

<210> 52
<211> 163
<212> PRT
<213> Homosapien

<400> 52
Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
1 5 10 15
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Gly

20	25	30
Ser Asn Asn Lys Ser Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln		
35	40	45
Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val		
50	55	60
Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr		
65	70	75
Ile Ser Ser Leu Gln Ala Ala Asp Val Ala Val Tyr Tyr Cys Gln Gln		
85	90	95
His Tyr Ser Thr Pro Cys Ser Phe Gly Gln Gly Thr Lys Leu Glu Ile		
100	105	110
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp		
115	120	125
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn		
130	135	140
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu		
145	150	155
Gln Ser Gly		160

<210> 53
 <211> 550
 <212> DNA
 <213> Homosapien

<400> 53
 caggtgcagc tgggtgcagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
 tcctgcagg cttctggata caccttcacc ggctactatc tgcactgggt gcgacaggcc 120
 cctggacaag ggcttgagtg gatgggatgg atcaaccctt acaatgatgg cacaactat 180
 gcacagaagt ttcaggcagc ggtcaccatg accaggacac cgtccatcag cacagcctac 240
 atggagctga gcaggctgag atctgacgac acggccgtt attactgtgc gagagatata 300
 gccgcagctg gagccgtcta ctttgactac tggggccagg gaaccctggt caccgtctcc 360
 tcagcttcca ccaagggccc atccgtcttc cccctggcgc cctgctccag gacacccctcc 420
 gagagcacag ccgcctggg ctgcctggc aaggactact ttcccccgaac cggtgacgg 480
 gtcgtggAAC tcaggcgcgg tgaccagcgg cgtgcacacc ttcccgctg tcctacagtc 540
 ctcaggactt 550

<210> 54
 <211> 183
 <212> PRT
 <213> Homosapien

<400> 54
 Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
 1 5 10 15
 Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Gly Tyr
 20 25 30
 Tyr Leu His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
 35 40 45
 Gly Trp Ile Asn Pro Tyr Asn Asp Gly Thr Asn Tyr Ala Gln Lys Phe
 50 55 60
 Gln Gly Arg Val Thr Met Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr
 65 70 75 80
 Met Glu Leu Ser Arg Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys

85	90	95
Ala Arg Asp Ile Ala Ala Ala Gly Ala Val Tyr Phe Asp Tyr Trp Gly		
100	105	110
Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser		
115	120	125
Val Phe Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala		
130	135	140
Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Arg Thr Gly Asp Gly		
145	150	155
Val Val Glu Leu Arg Arg Pro Asp Gln Arg Arg Ala His Leu Pro Gly		
165	170	175
Cys Pro Thr Val Leu Arg Thr		
180		

<210> 55
 <211> 458
 <212> DNA
 <213> Homosapien

<400> 55
 gacatccaga tgacccagtc tccatccctc ctgtctgcat ctgttaggaga cagagtcacc 60
 atcaattgcc aggcgagtca ggacattacc acctatttaa attggatca gcagaaacca 120
 gggaaagccc ctaagctctt gatctacat gcatccaatt tggaaacagg ggtcccatca 180
 aggttcagtg gaagtggatc tgggacagat ttactttca ccatcagcag cctgcagcct 240
 gaagatattt caacatatta ctgtcaacaa tatgataatc tcccgatcac cttcggccaa 300
 gggacacgac tggagattaa acgaactgtg gctgcaccat ctgtcttcat cttccggcca 360
 tctgatgagc agttgaaatc tggaaactgcc tctgttgtgt gcctgctgaa taacttctat 420
 cccagagagg ccaaagtaca gggaaagggtgg ataaacgcc 458

<210> 56
 <211> 152
 <212> PRT
 <213> Homosapien

<400> 56
 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
 1 5 10 15
 Asp Arg Val Thr Ile Thr Cys Gln Ala Ser Gln Asp Ile Thr Thr Tyr
 20 25 30
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
 35 40 45
 Tyr Asp Ala Ser Asn Leu Glu Thr Gly Val Pro Ser Arg Phe Ser Gly
 50 55 60
 Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Leu Gln Pro
 65 70 75 80
 Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Tyr Asp Asn Leu Pro Ile
 85 90 95
 Thr Phe Gly Gln Gly Thr Arg Leu Glu Ile Lys Arg Thr Val Ala Ala
 100 105 110
 Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly
 115 120 125
 Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala
 130 135 140
 Lys Val Gln Gly Arg Trp Ile Thr

145

150

<210> 57
<211> 571
<212> DNA
<213> Homosapien

<400> 57
caggtccagc tggcacagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcagg tttccggata caccctcaact gaattatcca tgcactgggt gcgacaggct 120
cctggaaaag ggcttgagtg gatgggaggt tttgatcctg aagatggtga aacaatctac 180
gcacagaagt tccagggcag agtcatgatg accgaggaca catctacaga cacagcctc 240
atggacctga gcagcctgag atctgaggac acggccgtgt attactgtgc aacagacgat 300
atgttgaccc ctcactaccc tctactcggt atggacgtct ggggccaagg gaccacggtc 360
accgtctccct cagcttccac caagggccca tccgtttcc ccctggcgcc ctgctccagg 420
agcacctccg agagcacacgc cgccctggc tgcttgcaggacta accagcggcg tgcacaccc 480
gtgacgggtgt cgtggaactc aggcgcctg accagcggcg tgcacaccc 540
ctacagtccct caggactcta ctccctcagc a 571

<210> 58
<211> 190
<212> PRT
<213> Homosapien

<400> 58
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Met Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Phe
65 70 75 80
Met Asp Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr Asp Asp Met Leu Thr Pro His Tyr Leu Tyr Phe Gly Met Asp
100 105 110
Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Ser Thr Lys
115 120 125
Gly Pro Ser Val Phe Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu
130 135 140
Ser Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro
145 150 155 160
Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr
165 170 175
Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser
180 185 190

<210> 59
<211> 458
<212> DNA

<213> Homosapien

<400> 59

gacatccaga tgacccagtc tccatccccc ctgtctgcat ctgttaggaga cagagtcacc 60
atcaacttgc gggcaagtca gggcattaga aatgatttag gctggtatca gcagaaacca 120
gggaaagccc ctaagcgcct gatctatgct acatccagtt tgcaaagtgg ggtccatca 180
aggttcagcg gcagtggatc tgggacagaa ttcaactctca caatcagcag cctgcagcct 240
gaagattttg caacttatta ctgtctacag cataatactt acccattcac tttcgccct 300
gggaccaaag tggatatcaa acgaactgtg gctgcaccat ctgtcttcat cttccgc 360
tctgatgagc agttgaaatc tggaaactgcc tctgttgtt gcctgtgaa taacttctat 420
cccagagagg ccaaagtaca gtggaaagggtg gataacgc 458

<210> 60

<211> 152

<212> PRT

<213> Homosapien

<400> 60

Asp	Ile	Gln	Met	Thr	Gln	Ser	Pro	Ser	Ser	Leu	Ser	Ala	Ser	Val	Gly
1										10					15
Asp	Arg	Val	Thr	Ile	Thr	Cys	Arg	Ala	Ser	Gln	Gly	Ile	Arg	Asn	Asp
				20						25					30
Leu	Gly	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	Lys	Ala	Pro	Lys	Arg	Leu	Ile
				35						40					45
Tyr	Ala	Thr	Ser	Ser	Leu	Gln	Ser	Gly	Val	Pro	Ser	Arg	Phe	Ser	Gly
				50						55					60
Ser	Gly	Ser	Gly	Thr	Glu	Phe	Thr	Leu	Thr	Ile	Ser	Ser	Leu	Gln	Pro
				65						70					80
Glu	Asp	Phe	Ala	Thr	Tyr	Tyr	Cys	Leu	Gln	His	Asn	Thr	Tyr	Pro	Phe
				85						90					95
Thr	Phe	Gly	Pro	Gly	Thr	Lys	Val	Asp	Ile	Lys	Arg	Thr	Val	Ala	Ala
				100					105						110
Pro	Ser	Val	Phe	Ile	Phe	Pro	Pro	Ser	Asp	Glu	Gln	Leu	Lys	Ser	Gly
				115					120						125
Thr	Ala	Ser	Val	Val	Cys	Leu	Leu	Asn	Asn	Phe	Tyr	Pro	Arg	Glu	Ala
				130					135						140
Lys	Val	Gln	Trp	Lys	Val	Asp	Asn								
				145					150						

<210> 61

<211> 1338

<212> DNA

<213> Homosapien

<400> 61

caggtgcagc tgcaggagtc gggcccgagga ctggtaagc cttcacagac cctgtccctc 60
acctgcactg tctcagggtgg ctccatcagc agtgggtggta actactggaa ctggatccgc 120
cagcacccag ggaaggccct ggagtggatt gggtacatct attacagtgg aaacacctac 180
tacaaccctgt ccctcaagag tcgaattacc atatcaatag acacgtctaa gaaccagttc 240
tccctgaccc tgagctctgt gactgcccgcg gacacggccg tgtattactg tgcgagagat 300
ggtggagacg atgctttga tatctgggc caagggacaa tggtcaccgt ctcttcagct 360
tccaccaagg gcccattccgt cttcccccgt gcgcctgtccctg ccaggagcac ctccgagagc 420
acagccgccc tgggctgcct ggtcaaggac tacttccccg aaccggtgac ggtgtcgtgg 480
aactcaggcg ccctgaccag cggcgtgcac accttccccgg ctgtcctaca gtcctcagga 540

ctctactccc tcagcagcgt ggtgaccgtg ccctccagca gcttggcac gaagacctac 600
 acctgcaacg tagatcacaa gcccagcaac accaagggtgg acaagagagt tgagtccaaa 660
 tatggtcccc catgcccata atgcccagca cctgagttcc tggggggacc atcagtcttc 720
 ctgttccccca caaaaacccaa ggacactctc atgatctccc ggaccctcga ggtcacgtgc 780
 gtgggtgggg acgtgagcca ggaagacccc gaggtccagt tcaactggta cgtggatggc 840
 gtggaggtgc ataatgccaa gacaaagccg cggaggagc agttcaacag cacgtaccgt 900
 gtggtcagcg tcctcaccgt cctgcaccag gactggctga acggcaagga gtacaagtgc 960
 aaggcttcca acaaaggcct cccgtctcc atcgagaaaa ccatctccaa agccaaagg 1020
 cagccccgag agccacaggt gtacaccctg ccccatccc aggaggagat gaccaagaac 1080
 caggtcagcc tgacctgcct ggtcaaaggc ttctacccca ggcacatcgc cgtggagtgg 1140
 gagagcaatg ggcagccgga gaacaactac aagaccacgc ctcccgtgct ggactccgac 1200
 ggctcttcttc tcctctacag caggctaacc gtggacaaga gcaggtggca ggagggaaat 1260
 gtcttctcat gctccgtat gcatgaggt ctgcacaacc actacacaca gaagagcctc 1320
 tccctgtctc tgggtaaa 1338

<210> 62
 <211> 446
 <212> PRT
 <213> Homosapien

<400> 62
 Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Lys Pro Ser Gln
 1 5 10 15
 Thr Leu Ser Leu Thr Cys Thr Val Ser Gly Gly Ser Ile Ser Ser Gly
 20 25 30
 Gly Asn Tyr Trp Asn Trp Ile Arg Gln His Pro Gly Lys Gly Leu Glu
 35 40 45
 Trp Ile Gly Tyr Ile Tyr Tyr Ser Gly Asn Thr Tyr Tyr Asn Pro Ser
 50 55 60
 Leu Lys Ser Arg Ile Thr Ile Ser Ile Asp Thr Ser Lys Asn Gln Phe
 65 70 75 80
 Ser Leu Thr Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr
 85 90 95
 Cys Ala Arg Asp Gly Gly Asp Asp Ala Phe Asp Ile Trp Gly Gln Gly
 100 105 110
 Thr Met Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
 115 120 125
 Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
 130 135 140
 Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
 145 150 155 160
 Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu
 165 170 175
 Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser
 180 185 190
 Ser Ser Leu Gly Thr Lys Thr Tyr Thr Cys Asn Val Asp His Lys Pro
 195 200 205
 Ser Asn Thr Lys Val Asp Lys Arg Val Glu Ser Lys Tyr Gly Pro Pro
 210 215 220
 Cys Pro Ser Cys Pro Ala Pro Glu Phe Leu Gly Pro Ser Val Phe
 225 230 235 240
 Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro
 245 250 255
 Glu Val Thr Cys Val Val Val Asp Val Ser Gln Glu Asp Pro Glu Val
 260 265 270

Gln Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr
 275 280 285
 Lys Pro Arg Glu Glu Gln Phe Asn Ser Thr Tyr Arg Val Val Ser Val
 290 295 300
 Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys
 305 310 315 320
 Lys Val Ser Asn Lys Gly Leu Pro Ser Ser Ile Glu Lys Thr Ile Ser
 325 330 335
 Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro
 340 345 350
 Ser Gln Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val
 355 360 365
 Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly
 370 375 380
 Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp
 385 390 395 400
 Gly Ser Phe Phe Leu Tyr Ser Arg Leu Thr Val Asp Lys Ser Arg Trp
 405 410 415
 Gln Glu Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His
 420 425 430
 Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Leu Gly Lys
 435 440 445

<210> 63
 <211> 642
 <212> DNA
 <213> Homosapien

<400> 63
 gacatccaga tgacccagtc tccatccctcc ctgtctgcat ctgttaggaga cagagtcacc 60
 atcaacttgcg aggccgagtca ggacattagc aactatttaa attggatcatca gcagaaaacca 120
 gggaaagccc ctaaaactcct gatctacat gcatccaatt tggaaacagg ggtcccatca 180
 aggttcagtg gaagtggatc tgggacagat tttactttca ccatcaacag cctgcagcct 240
 gaagatattt caacatatta ctgtcaagaa tataataatc tcccgtacag ttttggccag 300
 gggaccaagt tggagatcaa acgaactgtg gctgcaccat ctgtcttcat cttcccgcca 360
 tctgatgagc agttgaaatc tggaaactgccc tctgttgtgt gcctgctgaa taacttctat 420
 cccagagagg ccaaagtaca gtggaaagggtg gataacgccc tccaatcggg taactcccag 480
 gagagtgtca cagagcagga cagcaaggac agcacctaca gcctcagcag caccctgacg 540
 ctgagcaaag cagactacga gaaaacacaaa gtctacgcct gcgaagtac ccacatcaggc 600
 ctgagctcgc ccgtcacaaa gagcttcaac agggagagt gt 642

<210> 64
 <211> 214
 <212> PRT
 <213> Homosapien

<400> 64
 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
 1 5 10 15
 Asp Arg Val Thr Ile Thr Cys Gln Ala Ser Gln Asp Ile Ser Asn Tyr
 20 25 30
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
 35 40 45
 Tyr Asp Ala Ser Asn Leu Glu Thr Gly Val Pro Ser Arg Phe Ser Gly

50	55	60													
Ser	Gly	Ser	Gly	Thr	Asp	Phe	Thr	Phe	Thr	Ile	Asn	Ser	Leu	Gln	Pro
65															80
Glu	Asp	Ile	Ala	Thr	Tyr	Tyr	Cys	Gln	Glu	Tyr	Asn	Asn	Leu	Pro	Tyr
															95
Ser	Phe	Gly	Gln	Gly	Thr	Lys	Leu	Glu	Ile	Lys	Arg	Thr	Val	Ala	Ala
															100
Pro	Ser	Val	Phe	Ile	Phe	Pro	Pro	Ser	Asp	Glu	Gln	Leu	Lys	Ser	Gly
															105
Thr	Ala	Ser	Val	Val	Cys	Leu	Leu	Asn	Asn	Phe	Tyr	Pro	Arg	Glu	Ala
															110
Lys	Val	Gln	Trp	Lys	Val	Asp	Asn	Ala	Leu	Gln	Ser	Gly	Asn	Ser	Gln
															115
Glu	Ser	Val	Thr	Glu	Gln	Asp	Ser	Lys	Asp	Ser	Thr	Tyr	Ser	Leu	Ser
															120
Ser	Thr	Leu	Thr	Leu	Ser	Lys	Ala	Asp	Tyr	Glu	Lys	His	Lys	Val	Tyr
															125
Ala	Cys	Glu	Val	Thr	His	Gln	Gly	Leu	Ser	Ser	Pro	Val	Thr	Lys	Ser
															130
Phe	Asn	Arg	Gly	Glu	Cys										
															135
															140
															145
															150
															155
															160
															165
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															185
															190
															195
															200
															205
															210

<210> 65
 <211> 1341
 <212> DNA
 <213> Homosapien

<400> 65
 caggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgcaggtc 60
 tcctgcaagg tttccggaga caccctact gaattatcca tgcactgggt gcgacaggct 120
 cctggaaaag ggcttgagt gatggaggt tttgatcctg aagatggtga aacaatctac 180
 gcacggaaatg tccaggccag agtcaccatg accgaggaca catctacaga cacagttac 240
 atggagctga gcagcctgag atctgaggac acgcccgtgt atttctgtgc aacagattca 300
 cgtggatata gtggctactt tgacaactgg ggcaggaa ccctggtcac cgctcctca 360
 gcttccacca agggcccatc cgtcttcccc ctggccccc gctccaggag caccctccgag 420
 agcacagccg ccctgggctg cctggtcaag gactacttcc ccgaaccgg gacgggtgtcg 480
 tggaaactcag gcccctgac cagcggcgtg cacaccccttcc cggctgtccct acagtcctca 540
 ggactctact ccctcagcag cgtggtgacc gtggcccttca gcagcttggg cacaagacc 600
 tacacctgca acgttagatca caagcccagc aacaccaagg tggacaagag agttgagtcc 660
 aaatatggtc ccccatgccc atcatgccc gcacctgagt tcctgggggg accatcagtc 720
 ttcctgttcc ccccaaaaacc caaggacact ctcatgatct cccggacccc tgaggtcagc 780
 tgcgtggtgg tggacgtgag ccaggaagac cccgagggtcc agttcaactg gtacgtggat 840
 ggcgtggagg tgcataatgc caagacaaag cccggggagg agcagttcaa cagcacgtac 900
 cgtgtggtca gcgtccctcac cgtcctgcac cagactggc tgaacggcaa ggagtacaag 960
 tgcacgggtct ccaacaaaagg cctcccgccc tccatcgaga aaaccatctc caaagccaaa 1020
 gggcagccccc gagagccaca ggtgtacacc ctggccccc gccaggagga gatgaccaag 1080
 aaccaggtca gcctgacactg cctggtcaaa ggcttctacc ccagcgcacat cgcgggtgg 1140
 tgggagagca atgggcagcc ggagaacaa tacaagacca cgcctccctgt gctggactcc 1200
 gacggctcct tcttcctcta cagcaggcta accgtggaca agagcagggtg gcaaggaggg 1260
 aatgtcttct catgctccgt gatgcatgag gctctgcaca accactacac acagaagagc 1320
 ctctccctgt ctctggtaa a 1341

<210> 66
 <211> 447

<212> PRT
<213> Homosapien

<400> 66
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Gln Val Ser Cys Lys Val Ser Gly Asp Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Arg Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Val Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Phe Cys
85 90 95
Ala Thr Asp Ser Arg Gly Tyr Ser Gly Tyr Phe Asp Asn Trp Gly Gln
100 105 110
Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val
115 120 125
Phe Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala
130 135 140
Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser
145 150 155 160
Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val
165 170 175
Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro
180 185 190
Ser Ser Ser Leu Gly Thr Lys Thr Tyr Thr Cys Asn Val Asp His Lys
195 200 205
Pro Ser Asn Thr Lys Val Asp Lys Arg Val Glu Ser Lys Tyr Gly Pro
210 215 220
Pro Cys Pro Ser Cys Pro Ala Pro Glu Phe Leu Gly Gly Pro Ser Val
225 230 235 240
Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr
245 250 255
Pro Glu Val Thr Cys Val Val Val Asp Val Ser Gln Glu Asp Pro Glu
260 265 270
Val Gln Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys
275 280 285
Thr Lys Pro Arg Glu Glu Gln Phe Asn Ser Thr Tyr Arg Val Val Ser
290 295 300
Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys
305 310 315 320
Cys Lys Val Ser Asn Lys Gly Leu Pro Ser Ser Ile Glu Lys Thr Ile
325 330 335
Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro
340 345 350
Pro Ser Gln Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu
355 360 365
Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn
370 375 380
Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser
385 390 395 400
Asp Gly Ser Phe Phe Leu Tyr Ser Arg Leu Thr Val Asp Lys Ser Arg

405	410	415
Trp Gln Glu Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu		
420	425	430
His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Leu Gly Lys		
435	440	445

<210> 67
 <211> 660
 <212> DNA
 <213> Homosapien

<400> 67
 gacatcgta tgacccagtc tccagactcc ctggctgtgt ctctggcga gagggccacc 60
 atcaactgca agtccagcca gagtgtttta tacagctcca acaataacaa ctacttagtt 120
 tggtaccagc agaaaccagg acagcctcct aaattgctca tttactggc atctaccgg 180
 gaattcgggg ttccgtaccg attcagtgcc agccggctcg ggacagattt cactctcacc 240
 atcagcagcc tgcaggctga agatgtggca gtttattact gtcagcaata ttattttct 300
 ccgtggacgt tcggccaagg gaccaaggtg gaaatcaaac gaactgtggc tgcaccatct 360
 gtcttcatct tccccccatc tgcaggccatc ttgaaatctg gaactgcctc tgggtgtgc 420
 ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgccctc 480
 caatcgggta actcccagga gagtgtcaca gaggcaggaca gcaaggacag cacctacagc 540
 ctcagcagca ccctgacgct gagcaaagca gactacgaga aacacaagt ctacgcctgc 600
 gaagtcaccc atcaggccct gagtcgccc gtacacaaga gcttcaacag gggagagtgt 660

<210> 68
 <211> 220
 <212> PRT
 <213> Homosapien

<400> 68
 Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
 1 5 10 15
 Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
 20 25 30
 Ser Asn Asn Asn Asn Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln
 35 40 45
 Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Phe Gly Val
 50 55 60
 Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
 65 70 75 80
 Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
 85 90 95
 Tyr Tyr Phe Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
 100 105 110
 Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
 115 120 125
 Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
 130 135 140
 Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu
 145 150 155 160
 Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp
 165 170 175
 Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr

180	185	190
Glu Lys His Lys Val Tyr Ala Cys	Glu Val Thr His Gln Gly Leu Ser	
195	200	205
Ser Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys		
210	215	220

<210> 69
 <211> 556
 <212> DNA
 <213> Homosapien

<400> 69

caggtccagc tggcacagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
 tcctgcagg tttccggata caccctcaact gatttatcca tgcactgggt gcgacaggct 120
 cctggaaaag ggcttgagtg gatggggaggt tttgatcctg aagatggtga aacaatctac 180
 gcacagaagt tccaggcag agtcaccatg accgaggaca catcttcaga cacagcctac 240
 atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aaccacgaa 300
 ttttggagtg gttatattga ctactgggc cagggAACCC tggtcaccgt ctccctcagct 360
 tccaccaagg gcccattccgt cttcccccgt gcgcctgtgt ccaggagcac ctccgagagc 420
 acagccgccc tgggctgcct ggtcaaggac tacttccccc aaccgggtgac ggtgtcgtgg 480
 aactcaggcg ccctgaccag cggcgtgcac accttcccg ctgtcctaca gtcctcagga 540
 ctctactccc tcagca 556

<210> 70

<211> 185
 <212> PRT
 <213> Homosapien

<400> 70

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala			
1	5	10	15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Asp Leu			
20	25	30	
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met			
35	40	45	
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe			
50	55	60	
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Ser Asp Thr Ala Tyr			
65	70	75	80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys			
85	90	95	
Ala Thr His Glu Phe Trp Ser Gly Tyr Phe Asp Tyr Trp Gly Gln Gly			
100	105	110	
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe			
115	120	125	
Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu			
130	135	140	
Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp			
145	150	155	160
Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu			
165	170	175	
Gln Ser Ser Gly Leu Tyr Ser Leu Ser			
180	185		

<210> 71
<211> 476
<212> DNA
<213> Homosapien

<400> 71
gacatcgtga tgacccagtc tccagactcc ctggctgtgt ctctggcga gagggccacc 60
atcaactgca agtccagcca gagtgttta ttcaagctcca acaataagag ctacttaact 120
tggtaccagc agaaaccagg acagcctccct aaattactca ttttctggc atctatccgg 180
gaatccgggg tccctgaccg aatcagtgacgc agcgggtctg ggacagatct cactctcacc 240
atcagcagcc tgcaggctga agatgcgca gtttattact gtcagcaata ttatagtagt 300
ccgtggacgt tcggccaagg gaccaaggtg gaaatcaaac gaactgtggc tgcaccatct 360
gtcttcatct tccccccatc tgatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaaggtgga taacgc 476

<210> 72
<211> 158
<212> PRT
<213> Homosapien

<400> 72
Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
1 5 10 15
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Phe Ser
20 25 30
Ser Asn Asn Lys Ser Tyr Leu Thr Trp Tyr Gln Gln Lys Pro Gly Gln
35 40 45
Pro Pro Lys Leu Leu Ile Phe Trp Ala Ser Ile Arg Glu Ser Gly Val
50 55 60
Pro Asp Arg Ile Ser Gly Ser Gly Ser Gly Thr Asp Leu Thr Leu Thr
65 70 75 80
Ile Ser Ser Leu Gln Ala Glu Asp Ala Ala Val Tyr Tyr Cys Gln Gln
85 90 95
Tyr Tyr Ser Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
100 105 110
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
115 120 125
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
130 135 140
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn
145 150 155

<210> 73
<211> 546
<212> DNA
<213> Homosapien

<400> 73
caggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcaagg ttccggata caccctcactt gaattatcca tgcactgggt ggcacaggct 120
cctggaaaag ggcttgatgt gatgggaggt tttgatcctg aagatggta aataatccac 180
gcacagaagt tccagggcag agtcaccatg accgaggaca catctacaga cacagcctac 240
atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aacaggcgat 300

ttttggagtg gttattacct tgactggtgg ggcagggaa ccctggcac cgttcctca 360
gcttccacca agggccatc cgtttcccc ctggcgccct gctccaggag cacctccgag 420
agcacagccg ccctgggctg cctggtaag gactacttc ccgaaccggg gacgggtgtcg 480
tggaaacttag gcccctgac cagcggcgtg cacacccctcc cggctgtcct acagtccctca 540
ggactt 546

<210> 74
<211> 182
<212> PRT
<213> Homosapien

<400> 74
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Ser Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Ile Ile His Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr Gly Asp Phe Trp Ser Gly Tyr Tyr Leu Asp Trp Trp Gly Gln
100 105 110
Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val
115 120 125
Phe Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala
130 135 140
Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser
145 150 155 160
Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val
165 170 175
Leu Gln Ser Ser Gly Leu
180

<210> 75
<211> 457
<212> DNA
<213> Homosapien

<400> 75
gaaatagtga tgatgcagtc tccagccacc ctgtctgtgt ctccagggga aagagccacc 60
ctctcctgca gggccagtca gagtgttaac agcaacttag cctggcacca gcagaaacct 120
ggccaggctc ccaggctcct catcaacggt gcatccacca gggccactgg catcccagcc 180
aggttcagtg gcagtgggtc tgggacagag ttcaccctca ccatcagcag cctgcagttct 240
gaagattttt caatttatta ctgtcagcag tataatgact ggcctacgtt cactttcgcc 300
ggagggacca aggtggagat caatcgaact gtggctgcac catctgtctt catcttcccg 360
ccatctgatg agcagttgaa atctggaaact gcctctgttg tgcctgct gaataacttc 420
tatcccagag aggccaaagt acagtggaa ggtggat 457

<210> 76
<211> 152

<212> PRT
<213> Homosapien

<400> 76

Glu	Ile	Val	Met	Met	Gln	Ser	Pro	Ala	Thr	Leu	Ser	Val	Ser	Pro	Gly	
1					5					10					15	
Glu	Arg	Ala	Thr	Leu	Ser	Cys	Arg	Ala	Ser	Gln	Ser	Val	Asn	Ser	Asn	
										20			25		30	
Leu	Ala	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	Gln	Ala	Pro	Arg	Leu	Leu	Ile	
										35		40		45		
Asn	Gly	Ala	Ser	Thr	Arg	Ala	Thr	Gly	Ile	Pro	Ala	Arg	Phe	Ser	Gly	
										50		55		60		
Ser	Gly	Ser	Gly	Thr	Glu	Phe	Thr	Leu	Thr	Ile	Ser	Ser	Leu	Gln	Ser	
										65		70		75		80
Glu	Asp	Phe	Ala	Ile	Tyr	Tyr	Cys	Gln	Gln	Tyr	Asn	Asp	Trp	Pro	Thr	
										85		90		95		
Phe	Thr	Phe	Gly	Gly	Thr	Lys	Val	Glu	Ile	Asn	Arg	Thr	Val	Ala		
									100		105		110			
Ala	Pro	Ser	Val	Phe	Ile	Phe	Pro	Pro	Ser	Asp	Glu	Gln	Leu	Lys	Ser	
									115		120		125			
Gly	Thr	Ala	Ser	Val	Val	Cys	Leu	Leu	Asn	Asn	Phe	Tyr	Pro	Arg	Glu	
									130		135		140			
Ala	Lys	Val	Gln	Trp	Glu	Gly	Gly									
					145		150									

<210> 77
<211> 470
<212> DNA
<213> Homosapien

<400> 77

caggtccagc	tgg tac agtc	tggggctgag	gtgaagaagc	ctggggcctc	agtgaaggtc	60
tcctgcaagg	tttccggata	caccctcaact	gaattatcca	tgcactgggt	gcgacaggct	120
cctggaaaag	ggctttagtg	gatgggaggt	tttgatcctg	aagatggtga	aacaatgtac	180
gcacagaagt	tccaggcag	agt caccatg	accaggaca	catctacaga	cacagcctac	240
atggagctga	gcagcctgag	atctgaggac	acggccgtgt	attactgtgc	aaccgacgat	300
ttttggagtg	gttattttga	ctactggggc	cagggAACCC	tggtcaccgt	ctcctcagcc	360
tccaccaagg	gcccattcggt	cttccccctg	gcgcctgtct	ccaggagcac	ctccgagagc	420
acagcggccc	tggctgcct	ggtcaaggac	tacttccccg	aaccggcagg		470

<210> 78
<211> 156
<212> PRT
<213> Homosapien

<400> 78

Gln	Val	Gln	Leu	Val	Gln	Ser	Gly	Ala	Glu	Val	Lys	Lys	Pro	Gly	Ala
1					5					10					15
Ser	Val	Lys	Val	Ser	Cys	Lys	Val	Ser	Gly	Tyr	Thr	Leu	Thr	Glu	Leu
										20		25		30	
Ser	Met	His	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Met
										35		40		45	
Gly	Gly	Phe	Asp	Pro	Glu	Asp	Gly	Glu	Thr	Met	Tyr	Ala	Gln	Lys	Phe
										50		55		60	

Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
 65 70 75 80
 Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Thr Asp Asp Phe Trp Ser Gly Tyr Phe Asp Tyr Trp Gly Gln Gly
 100 105 110
 Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
 115 120 125
 Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
 130 135 140
 Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Ala
 145 150 155

<210> 79
 <211> 490
 <212> DNA
 <213> Homosapien

<400> 79
 gacatcgta tgacccagtc tccagactcc ctggctgtgt ctctggacga gagggccacc 60
 atcaactgca agtccagcca gagtgttta tacagtccca accaaaagaa ctacttagtt 120
 tggtatcagc agaagccagg acagcctcct aagctgctcc ttactgggc atctatccgg 180
 gaatccgggg tccctgaccg attcagtgcc agccgggtctg ggacagattt cactctcacc 240
 atcagcagcc tgcaggctga agatgtggca gtttattact gtcaacaaag ttatttact 300
 ccgtggacgt tcggccaagg gaccaagggtg gaaatcaaac gaactgtggc tgccaccatct 360
 gtcttcatct tcccggccatc tcatgtggccatc ttgaaaatctg gaactgcctc tgggtgtgc 420
 ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacggccctc 480
 caatcggtta 490

<210> 80
 <211> 163
 <212> PRT
 <213> Homosapien

<400> 80
 Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Asp
 1 5 10 15
 Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
 20 25 30
 Pro Asn Gln Lys Asn Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln
 35 40 45
 Pro Pro Lys Leu Leu Tyr Trp Ala Ser Ile Arg Glu Ser Gly Val
 50 55 60
 Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
 65 70 75 80
 Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
 85 90 95
 Ser Tyr Phe Thr Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
 100 105 110
 Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
 115 120 125
 Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
 130 135 140
 Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu

145
Gln Ser Gly

150

155

160

<210> 81
<211> 556
<212> DNA
<213> Homosapien

<400> 81
caggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcaagg tttccggata caccctcagt gaattatcca tgcactgggt gcgacaggct 120
cctggaaaag ggctttagtg gatgggaggt tttgatcctg aagatgtat aacaatctac 180
gcacagaagt tccagggcag agtcaccatg accgaggaca catctacaga cacagccttc 240
atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aaccacgat 300
tttggagtg gttatccatca ctactggggc cagggaaaccc tggtcaccgt ctccctagct 360
tccaccaagg gccccatccgt cttcccccgt gcgcctgct ccaggagcac ctccgagagc 420
acagccgccc tgggctgcct ggtcaaggac tacttcccg aaccgggtgac ggtgtcgtgg 480
aactcaggcg ccctgaccag cggcgtgcac accttcccg ctgtcctaca gtccctagga 540
ctctactcccc tcagca 556

<210> 82
<211> 185
<212> PRT
<213> Homosapien

<400> 82
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Ser Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Asp Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Phe
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr His Asp Phe Trp Ser Gly Tyr Phe His Tyr Trp Gly Gln Gly
100 105 110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125
Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
130 135 140
Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
145 150 155 160
Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu
165 170 175
Gln Ser Ser Gly Leu Tyr Ser Leu Ser
180 185

<210> 83

<211> 476
<212> DNA
<213> Homosapien

<400> 83
gacatcgtga tgacccagtc tccagactcc ctggctgtgt ctctggcga gaggggcacc 60
atcaactgca agtccagcca gagtgttta tacagctccg acaataagag ctacttagtt 120
tggtaccagc agaaaccagg acagcctccct aaggtgctca tttactgggc atctattcgg 180
gaatccgggg tccctgaccg attcagtgcc agccgggtctg ggacagattt cactctcacc 240
atcagcagcc tgcaggctga agatgtggca gtttattact gtcagcaata ttatactagt 300
ccgtggacgt tcggccaagg gaccaaggtg gaaatcaaac gaactgtggc tgcaccatct 360
gtcttcatct tcccgcacatc tgatgaggcag ttgaaatctg gaactgcctc tgggtgtgc 420
ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgc 476

<210> 84
<211> 158
<212> PRT
<213> Homosapien

<400> 84
Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
1 5 10 15
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
20 25 30
Ser Asp Asn Lys Ser Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln
35 40 45
Pro Pro Lys Val Leu Ile Tyr Trp Ala Ser Ile Arg Glu Ser Gly Val
50 55 60
Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65 70 75 80
Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
85 90 95
Tyr Tyr Thr Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
100 105 110
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
115 120 125
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
130 135 140
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn
145 150 155

<210> 85
<211> 543
<212> DNA
<213> Homosapien

<400> 85
caggtccagc tggtagcgtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgttaagg ttcccgata caccctcaact gaattatcca tgcactgggt gcgacaggct 120
cctggaaaag ggcttgatgt gatgggaggt tttatccctg aagatggta aacaatctac 180
gcacagaagt tccagggcag agtcaccatg accgaggaca catctacaga cacagcctac 240
atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aatccacgag 300
ttttggagtg gttatccatc ctactggggc cagggaaaccc tggtcaccgt ctccatcgatc 360
tccaccaagg gcccattccgt cttcccccgt gcgcctgtgt ccaggagcac ctccgagagc 420

acagccgccc tgggctgcct ggtcaaggac tacttccccg aaccgggtgac ggtgtcgtgg 480
aactcaggcg ccctgaccag cggcgtgcac accttcccggtgtcctaca gtcctcagga 540
ctt 543

<210> 86
<211> 181
<212> PRT
<213> Homosapien

<400> 86
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Ile His Glu Phe Trp Ser Gly Tyr Phe Asp Tyr Trp Gly Gln Gly
100 105 110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125
Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
130 135 140
Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
145 150 155 160
Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu
165 170 175
Gln Ser Ser Gly Leu
180

<210> 87
<211> 477
<212> DNA
<213> Homosapien

<400> 87
gacatcgta tgacccagtc tccagactcc ctggctgtgt ctctggcga gagggccacc 60
atcaactgca agtccagcct gagtgttta tacagctcca acaataagaa ctattttagtt 120
tggtaacctc agaaaccagg acagcctcct aagttgctca tttactggc atctacccgg 180
gaatccgggg tccctgaccg attcagtggc agcgggtctg ggacagatt cactctcacc 240
atcagcagcc tgcaggccga agatgtggca gtttattact gtcagcaata ttatagttct 300
ccgtggacgt tcggccaagg gaccaaggtg gaaatcaaac gaactgtggc tgcaccatct 360
gtcttcatct tccccccatc tgatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgcc 477

<210> 88
<211> 159
<212> PRT
<213> Homosapien

<400> 88

Asp	Ile	Val	Met	Thr	Gln	Ser	Pro	Asp	Ser	Leu	Ala	Val	Ser	Leu	Gly
1			5				10						15		
Glu	Arg	Ala	Thr	Ile	Asn	Cys	Lys	Ser	Ser	Leu	Ser	Val	Leu	Tyr	Ser
			20				25						30		
Ser	Asn	Asn	Lys	Asn	Tyr	Leu	Val	Trp	Tyr	Leu	Gln	Lys	Pro	Gly	Gln
			35				40					45			
Pro	Pro	Lys	Leu	Leu	Ile	Tyr	Trp	Ala	Ser	Thr	Arg	Glu	Ser	Gly	Val
			50				55				60				
Pro	Asp	Arg	Phe	Ser	Gly	Ser	Gly	Ser	Gly	Thr	Asp	Phe	Thr	Leu	Thr
	65			70				75					80		
Ile	Ser	Ser	Leu	Gln	Ala	Glu	Asp	Val	Ala	Val	Tyr	Tyr	Cys	Gln	Gln
			85				90					95			
Tyr	Tyr	Ser	Ser	Pro	Trp	Thr	Phe	Gly	Gln	Gly	Thr	Lys	Val	Glu	Ile
			100				105					110			
Lys	Arg	Thr	Val	Ala	Ala	Pro	Ser	Val	Phe	Ile	Phe	Pro	Pro	Ser	Asp
	115					120						125			
Glu	Gln	Leu	Lys	Ser	Gly	Thr	Ala	Ser	Val	Val	Cys	Leu	Leu	Asn	Asn
	130				135						140				
Phe	Tyr	Pro	Arg	Glu	Ala	Lys	Val	Gln	Trp	Lys	Val	Asp	Asn	Ala	
	145				150						155				

<210> 89

<211> 1335

<212> DNA

<213> Homosapien

<400> 89

caggtccagc	tggtagcagtc	tggggctgag	gtgaagaagc	ctggggcctc	agtgaaggtc	60
tcctgcagg	tttccggata	caccctcaact	gaattatcca	tgcactgggt	gcgacagact	120
cctggaaaag	ggcttgagtg	gatggggaggt	tttgcacttg	aagatggtga	aacaatctac	180
gcacagaagt	tccaggacag	agtcaccatg	accgaggaca	catctacaga	cacagcctac	240
atggaaactga	gcagcctgag	atctgaggac	acggccgtgt	attactgtgc	aacaaacgat	300
ttttggactg	gttattatga	ctactggggc	cagggAACCC	tggtcaccgt	ctcctcagcc	360
tccaccaagg	gcccattcggt	cttccccctg	gcgcctgtct	ccaggagcac	ctccgagagc	420
acagcggccc	tgggctgcct	ggtcaaggac	tacttccccg	aaccgggtac	ggtgtcggtgg	480
aactcaggcg	ctctgaccag	cggcgtgcac	accttcccag	ctgtcctaca	gtcctcagga	540
ctctactccc	tcagcagcgt	ggtgaccgtg	ccctccagca	acttcggcac	ccagacccatc	600
acctgcaacg	tagatcacaa	gcccagcaac	accaagggtgg	acaagacagt	tgagcgcaaa	660
tgttgtgtcg	agtggccacc	gtgcccagca	ccacctgtgg	caggaccgtc	agtcttcctc	720
ttccccccaa	aacccaagga	caccctcatg	atctcccgga	ccctcgagggt	cacgtgcgtg	780
gtgggtggacg	tgagccacga	agaccccgag	gtccagttca	actggtaagt	ggacggcgtg	840
gaggtgcata	atgccaagac	aaagccacgg	gaggagcagt	tcaacagcac	gttccgtgtg	900
gtcagegtcc	tcaccgttgc	gcaccaggac	tggctgaacg	gcaaggagta	caagtgcag	960
gtctccaaca	aaggcctccc	agccccatc	gagaaaaacca	tctccaaaac	caaagggcag	1020
ccccgagaac	cacagggtgt	caccctgccc	ccatccccggg	aggagatgac	caagaaccag	1080
gtcagcctga	cctgcctgtt	caaaggcttc	taccccaagcg	acatcgccgt	ggagtggag	1140
agcaatgggc	agccggagaa	caactacaag	accacacctc	ccatgctgga	ctccgacggc	1200
tccttcttcc	tctacagcaa	gctcaccgtg	gacaagagca	ggtggcagca	gggaaacgtc	1260
ttctctatgt	ccgtgtatgca	tgaggctctg	cacaaccact	acacgcagaa	gaggcctctcc	1320
ctgtctccgg	gtaaaa					1335

<210> 90

<211> 445
<212> PRT
<213> Homosapien

<400> 90
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Thr Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Asp Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr Asn Asp Phe Trp Thr Gly Tyr Asp Tyr Trp Gly Gln Gly
100 105 110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125
Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
130 135 140
Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
145 150 155 160
Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu
165 170 175
Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser
180 185 190
Ser Asn Phe Gly Thr Gln Thr Tyr Thr Cys Asn Val Asp His Lys Pro
195 200 205
Ser Asn Thr Lys Val Asp Lys Thr Val Glu Arg Lys Cys Cys Val Glu
210 215 220
Cys Pro Pro Cys Pro Ala Pro Pro Val Ala Gly Pro Ser Val Phe Leu
225 230 235 240
Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu
245 250 255
Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Gln
260 265 270
Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys
275 280 285
Pro Arg Glu Glu Gln Phe Asn Ser Thr Phe Arg Val Val Ser Val Leu
290 295 300
Thr Val Val His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys
305 310 315 320
Val Ser Asn Lys Gly Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys
325 330 335
Thr Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser
340 345 350
Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys
355 360 365
Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln
370 375 380
Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Met Leu Asp Ser Asp Gly
385 390 395 400

Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln
 405 410 415
 Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn
 420 425 430
 His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
 435 440 445

<210> 91
 <211> 660
 <212> DNA
 <213> Homosapien

<400> 91
 gacatcgta tgacccagtc tccagactcc ctggctgtgt ctctggcga gagggccacc 60
 atcaactgca agtccagcca gagtgttta tacagctcca acaataagaa ctacttagtt 120
 tggtaccaggc agaaaccagg acagcctctt aagacgctca ttactggc atctaccgg 180
 gaatccgggg tccctgaccg attcagtgcc agcgggtctg ggacagattt cactctcacc 240
 atcagcagcc tgcaggctga agatgtggg gtttattact gtcaacaata ttatactagt 300
 ccgtggacgt tcggccaagg gaccaaggtg gaaatcaagc gaactgtggc tgcaccatct 360
 gtcttcatct tcccgccatc tggatgacag ttgaaatctg gaactgcctc tggatgtgc 420
 ctgctgaata acttctatcc cagagagcc aaagtacagt ggaagggtgga taacgcctc 480
 caatcgggta actcccagga gagtgcaca gagcaggaca gcaaggacag cacctacagc 540
 ctcagcagca ccctgacgct gagcaaaagca gactacgaga aacacaaagt ctacgcctgc 600
 gaagtcaccc atcagggct gagctcgccc gtcacaaaga gcttcaacag gggagagtgt 660

<210> 92
 <211> 220
 <212> PRT
 <213> Homosapien

<400> 92
 Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
 1 5 10 15
 Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
 20 25 30
 Ser Asn Asn Lys Asn Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln
 35 40 45
 Pro Pro Lys Thr Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
 50 55 60
 Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
 65 70 75 80
 Ile Ser Ser Leu Gln Ala Glu Asp Val Gly Val Tyr Tyr Cys Gln Gln
 85 90 95
 Tyr Tyr Thr Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
 100 105 110
 Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
 115 120 125
 Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
 130 135 140
 Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu
 145 150 155 160
 Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp
 165 170 175

Ser	Thr	Tyr	Ser	Leu	Ser	Ser	Thr	Leu	Thr	Leu	Ser	Lys	Ala	Asp	Tyr
							180					185			190
Glu	Lys	His	Lys	Val	Tyr	Ala	Cys	Glu	Val	Thr	His	Gln	Gly	Leu	Ser
							195					200			205
Ser	Pro	Val	Thr	Lys	Ser	Phe	Asn	Arg	Gly	Glu	Cys				
						210			215						220

<210> 93
<211> 560
<212> DNA
<213> Homosapien

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<400> 93
caggtgcagc tgcaggagtc gggcccagga ctggtaaagc cgtcacagac cctgtccctc 60
acctgcactg tctctgggtgg ctccatcagc agtgggtggtt actactggag ctggatccgc 120
cagcacccag ggaaggggcct ggagtggatt ggttacatct attacagtgg gagcacctac 180
tacaaccctgt ccctcaagag tcgagttatc atatcgttag acacgtctaa gaaccagttc 240
tccctgaagc tgacctctgt gactgcccgcg gacacggcccg tgtattactg tgcgagatca 300
tatagcagct cgtccccact ggttcgaccc ctggggccag ggaaccctgg tcaccgtctc 360
ctcagcttcc accaaggggcc catccgtctt cccccctggcg ccctgtctca ggagcacctc 420
cgagagcaca gccccttgg gctgcctggt caaggactac ttccccgaac cggtgacgg 480
gtcgtggAAC tcaggcgccc tgaccagcgg cgtgcacacc ttcccggtg tcctacagtc 540
ctcaggactc tactccctca 560
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<210> 94
<211> 186
<212> PRT
<213> Homosapien

<400> 94
 Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Lys Pro Ser Gln
 1 5 10 15
 Thr Leu Ser Leu Thr Cys Thr Val Ser Gly Gly Ser Ile Ser Ser Gly
 20 25 30
 Gly Tyr Tyr Trp Ser Trp Ile Arg Gln His Pro Gly Lys Gly Leu Glu
 35 40 45
 Trp Ile Gly Tyr Ile Tyr Tyr Ser Gly Ser Thr Tyr Tyr Asn Pro Ser
 50 55 60
 Leu Lys Ser Arg Val Ile Ile Ser Val Asp Thr Ser Lys Asn Gln Phe
 65 70 75 80
 Ser Leu Lys Leu Thr Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr
 85 90 95
 Cys Ala Arg Ser Tyr Ser Ser Ser Pro Leu Val Arg Pro Leu Gly
 100 105 110
 Pro Gly Asn Pro Gly His Arg Leu Leu Ser Phe His Gln Gly Pro Ile
 115 120 125
 Arg Leu Pro Pro Gly Ala Leu Leu Gln Glu His Leu Arg Glu His Ser
 130 135 140
 Arg Pro Gly Leu Pro Gly Gln Gly Leu Leu Pro Arg Thr Gly Asp Gly
 145 150 155 160
 Val Val Glu Leu Arg Arg Pro Asp Gln Arg Arg Ala His Leu Pro Gly
 165 170 175
 Cys Pro Thr Val Leu Arg Thr Leu Leu Pro
 180 185

<210> 95
<211> 458
<212> DNA
<213> Homosapien

<400> 95
gacatccaga tgacccagtc tccatccccc ctgtctgcat ctgttaggaga cagagtcacc 60
atcacttgcc gggcaagtca gggcattaga aatgatttag gctggtatca gcagaaacca 120
gggaaagccc ctaagcgcct gatctatgt gcatccagtt tgcaaagtgg ggtcccatca 180
aggttcagcg gcagtggatc tgggacagaa ttcaactctca caatcagcag cctgcagcct 240
gaagattttg caacttatta ctgtctacag cataatagtt acccattcac tttcggccct 300
gggaccaaag tggatataaa acgaactgtg gctgcaccat ctgtcttcat cttcccgcca 360
tctgtatgagc agttgaaatc tggaaactgtc tctgtgtgt gcctgctgaa taacttctat 420
cccagagagg ccaaagtaca gtggaaaggta gataacgc 458

<210> 96
<211> 152
<212> PRT
<213> Homosapien

<400> 96
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1 5 10 15
Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Arg Asn Asp
20 25 30
Leu Gly Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Arg Leu Ile
35 40 45
Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
50 55 60
Ser Gly Ser Gly Thr Glu Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
65 70 75 80
Glu Asp Phe Ala Thr Tyr Tyr Cys Leu Gln His Asn Ser Tyr Pro Phe
85 90 95
Thr Phe Gly Pro Gly Thr Lys Val Asp Ile Lys Arg Thr Val Ala Ala
100 105 110
Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly
115 120 125
Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala
130 135 140
Lys Val Gln Trp Lys Val Asp Asn
145 150

<210> 97
<211> 559
<212> DNA
<213> Homosapien

<400> 97
caggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcaagg tttccggata caccctact gaattatcca tgcactgggt gcgacaggct 120
cctggaaaag ggcttgatgt gatgggaggt tttgatcctg aagatggta aacaatctac 180
gcacagaagt tccaggcag agtcaccatg accgaggaca catctacaga cacagcctac 240

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atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aacagatcgc 300
gagttttgga gtggttattt ctaccactgg ggccaggaa ccctggtcac cgtctcctca 360
gcctccacca agggcccatc ggtcttcccc ctggcgccct gctccaggag cacctccgag 420
agcacagcgg ccctgggctg cctggtcaag gactacttcc ccgaaccggg gacggtgtcg 480
tggaactctag gcgctctgac cagcggcgtg cacaccttcc cagctgtcct acagtccctca 540
ggactctact ccctcagca 559
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<210> 98
<211> 186
<212> PRT
<213> Homosapien

<400> 98

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
 1 5 10 15
 Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
 20 25 30
 Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
 35 40 45
 Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
 50 55 60
 Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
 65 70 75 80
 Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Thr Asp Arg Glu Phe Trp Ser Gly Tyr Phe Tyr His Trp Gly Gln
 100 105 110
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val
 115 120 125
 Phe Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala
 130 135 140
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser
 145 150 155 160
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val
 165 170 175
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser
 180 185

<210> 99
<211> 491
<212> DNA
<213> Homosapien

<400> 99

gacatcgta tgacccagtc tccagactcc ctggctgtgt ctctgggcga gagggccacc 60
atcaactgca agtccagcca gagtgttta tacagctcca acaatgagaa ctcttagct 120
tggtaccagc agaaaaccagg acagcctcct aaactgctca ttactggc atctaccgg 180
gaatccgggg tcccgagaccg cttcagtgcc agcgggtctg ggacagattt cactctcacc 240
atcagcagcc tgcaggctga agatgtggca gtttattact gtcagcaata ttataatagt 300
ccgtggacgt tcggccaagg gaccaaggtg gaaatcaaac gaactgtggc tgcaccatct 360
gtcttcatct tcccgccatc tgatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgcctcc 480
ccaatcggtt a 491

<210> 100
<211> 163
<212> PRT
<213> Homosapien

<400> 100
Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
1 5 10 15
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
20 25 30
Ser Asn Asn Glu Asn Phe Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln
35 40 45
Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
50 55 60
Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65 70 75 80
Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
85 90 95
Tyr Tyr Asn Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
100 105 110
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
115 120 125
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
130 135 140
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Ser
145 150 155 160
Pro Ile Gly

<210> 101
<211> 543
<212> DNA
<213> Homosapien

<400> 101
caggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcaagg tttccggata caccctcaact gaattatcca tgcactgggt gcgacaggct 120
cctggaaaag ggcttgagtg gatggggaggt tttgatcctg aagatggtga aacaatctac 180
gcacagaagt tccagggcag agtcaccatg accaggacatcatacagacac 240
atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aacggacgat 300
ttttggagtg gttatgttga ctactggggc cagggAACCC tggtcacccgt ctccctcagcc 360
tccaccaagg gccccatcggt ctccccctg gcgcctgtct ccaggagcac ctcccgagac 420
acagcggccc tgggctgcct ggtcaaggac tacttccccg aaccggtgac ggtgtcgtgg 480
aactcaggcg ctctgaccag cggcgtgcac accttcccg ctgtcctaca gtctcagga 540
ctt 543

<210> 102
<211> 181
<212> PRT
<213> Homosapien

<400> 102
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15

Ser	Val	Lys	Val	Ser	Cys	Lys	Val	Ser	Gly	Tyr	Thr	Leu	Thr	Glu	Leu
			20					25							30
Ser	Met	His	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Met
								35		40					45
Gly	Gly	Phe	Asp	Pro	Glu	Asp	Gly	Glu	Thr	Ile	Tyr	Ala	Gln	Lys	Phe
								50		55					60
Gln	Gly	Arg	Val	Thr	Met	Thr	Glu	Asp	Thr	Ser	Thr	Asp	Thr	Ala	Tyr
								65		70					80
Met	Glu	Leu	Ser	Ser	Leu	Arg	Ser	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys
								85		90					95
Ala	Thr	Asp	Asp	Phe	Trp	Ser	Gly	Tyr	Phe	Asp	Tyr	Trp	Gly	Gln	Gly
								100		105					110
Thr	Leu	Val	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe
								115		120					125
Pro	Leu	Ala	Pro	Cys	Ser	Arg	Ser	Thr	Ser	Glu	Ser	Thr	Ala	Ala	Leu
								130		135					140
Gly	Cys	Leu	Val	Lys	Asp	Tyr	Phe	Pro	Glu	Pro	Val	Thr	Val	Ser	Trp
								145		150					160
Asn	Ser	Gly	Ala	Leu	Thr	Ser	Gly	Val	His	Thr	Phe	Pro	Ala	Val	Leu
								165		170					175
Gln	Ser	Ser	Gly	Leu											
								180							

<210> 103
<211> 491
<212> DNA
<213> Homosapien

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<400> 103
gacatcgta tgacccagtc tccagactcc ctggctgtgt ctctggcga gaggccacc 60
atcaactgca agtccagtca gaggtttta tacaggtcta acaataagag ctacttagtt 120
tggtaccagc agaaactagg acagtctcct aagctgctca tttactggc atctacccgg 180
gaatccgggg tccctgaccg attcagtggc agcgggtctg ggacagattt cactctcacc 240
atcagcagcc tgcaggctga agatgtggca gtttattatt gtcaacaata ttatagttact 300
ccgtggacgt tcggccaaagg gaccaaggtg gaaatcaaacc gaaactgtggc tgcaccatct 360
gtcttcatct tcccgccatc tgatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgcctc 480
ccaatcggtt a 491
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<210> 104
<211> 163
<212> PRT
<213> Homosapien

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<400> 104
Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
  1           5           10          15
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Arg
  20          25          30
Ser Asn Asn Lys Ser Tyr Leu Val Trp Tyr Gln Gln Lys Leu Gly Gln
  35          40          45
Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
  50          55          60
Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr

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65	70	75	80
Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln			
85	90	95	
Tyr Tyr Ser Thr Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile			
100	105	110	
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp			
115	120	125	
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn			
130	135	140	
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu			
145	150	155	160
Pro Ile Gly			

<210> 105
 <211> 499
 <212> DNA
 <213> Homosapien

<400> 105
 cagggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
 tcctgcaagg ttccggata caccctact gaattatcca tgcactgggt ggcacaggct 120
 cctggaaaag ggcttgagtg gatgggaggt ttgatcctg aagatggtga aacaatctac 180
 gcacagaagt tccaggcgag agtcaccatg accgaggaca catctacaga cacagcctac 240
 atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aacagacgt 300
 ttttggagtg gttatgttga ctactgggc cagggAACCC tggtcaccgt ctccctagcc 360
 tccaccaagg gcccattcggt ctccccctg gcgcctgtt ccaggagcac ctccgagagc 420
 acagcggccc tgggctgcct ggtcaaggac tacttccccg aaccggtgac ggtgtcgtgg 480
 aactcaggcg ctctgacca 499

<210> 106
 <211> 166
 <212> PRT
 <213> Homosapien

<400> 106
 Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
 1 5 10 15
 Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
 20 25 30
 Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
 35 40 45
 Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
 50 55 60
 Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
 65 70 75 80
 Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Thr Asp Asp Phe Trp Ser Gly Tyr Phe Asp Tyr Trp Gly Gln Gly
 100 105 110
 Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
 115 120 125
 Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
 130 135 140

Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
145 150 155 160
Asn Ser Gly Ala Leu Thr
165

<210> 107
<211> 448
<212> DNA
<213> Homosapien

<400> 107
gacatcgta tgacccagtc tccagactcc ctggctgtgt ctctggcga gagggccacc 60
atcaactgca agtccagcca gagtgtttt acaataagaa ctacttagtt 120
tggtaccagc agaaaccagg acagcctcct aagctgctca ttactggc atctacccgg 180
gaatccgggg tccctgaccg attcagtgcc agcgggtctg ggacagattt cactctcacc 240
atcagcagcc tgcaaggctga agatgtggca gtttattact gtcagcaata ttatagtcct 300
acgtggacgt tcgccaagg gaccaaggtg gaaatcaaac gaactgtggc tgcaccatct 360
gtcttcatct tccgcctatc tgatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
ctgctgaata acttctatcc cagagagg 448

<210> 108
<211> 149
<212> PRT
<213> Homosapien

<400> 108
Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
1 5 10 15
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
20 25 30
Ser Asn Asn Lys Asn Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln
35 40 45
Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
50 55 60
Pro Asp Arg Phe Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65 70 75 80
Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
85 90 95
Tyr Tyr Ser Pro Thr Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
100 105 110
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
115 120 125
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
130 135 140
Phe Tyr Pro Arg Glu
145

<210> 109
<211> 540
<212> DNA
<213> Homosapien

<400> 109

caggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcaagg tttccggata caccctcaact gaattatcca tgcactgggt gcgacaggct 120
cctggaaaag ggctttagtg gatgggaggt tttgatcctg aagatggta aacaatctac 180
gcacagaagt tccagggcag agtcaccatg accgaggaca catctacaga cacagcctac 240
atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aacggacgat 300
ttttggagtg gttatggta ctactgggc cagggAACCC tggtagccgt ctccctcagcc 360
tccaccaagg gcccattcggt cttccctg gcgcctgct ccaggagcac ctccgagagc 420
acagcggccc tggctgcct ggtcaaggac tacttccccg aaccggtgac ggtgtcgtgg 480
aactcaggcg ctctgaccag cggcgtgcac accttcccag ctgtcctaca gtcctcagga 540

<210> 110
<211> 180
<212> PRT
<213> Homosapien

<400> 110
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr Asp Asp Phe Trp Ser Gly Tyr Phe Asp Tyr Trp Gly Gln Gly
100 105 110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125
Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
130 135 140
Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
145 150 155 160
Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu
165 170 175
Gln Ser Ser Gly
180

<210> 111
<211> 478
<212> DNA
<213> Homosapien

<400> 111
gacatcgta tgacccagtc tccagactcc ctggctgtgt ctctggcga gagggccacc 60
atcaactgca agtccagcca gagtgttttacagctcca acaataagaa ctacttagct 120
tggtagccagg agaaaccagg acagcctcct aagctgctca tttactggac atctacccgg 180
gaatccgggg tccctgaccg attcagtgcc agcgggtctg tgacagatt cactctcacc 240
atcagcagcc tgcaggctga agatgtggca gtttattact gtcagcaata ttatagttct 300
ccgtggacgt tcggccaagg gaccaagggtg gaaatcaaac gaactgtggc tgcaccatct 360

gtcttcatct tccggccatc tgatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgcct 478

<210> 112
<211> 159
<212> PRT
<213> Homosapien

<400> 112
Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
1 5 10 15
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
20 25 30
Ser Asn Asn Lys Asn Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln
35 40 45
Pro Pro Lys Leu Leu Ile Tyr Trp Thr Ser Thr Arg Glu Ser Gly Val
50 55 60
Pro Asp Arg Phe Ser Gly Ser Gly Ser Val Thr Asp Phe Thr Leu Thr
65 70 75 80
Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
85 90 95
Tyr Tyr Ser Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
100 105 110
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
115 120 125
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
130 135 140
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala
145 150 155

<210> 113
<211> 542
<212> DNA
<213> Homosapien

<400> 113
caggtccagc tggtagcgtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
tcctgcaagg ttccggata caccctcagt gaattatcca tgcactgggt gcacaggct 120
cctggaaaag ggctttagtg gatgggaggt ttgatcctg aagatggta aacaatctac 180
gcacagaagt tccaggccag agtcaccatg accaggacatcatacaca cacagcctac 240
atggagctga gcagcctgag atctgaggac acggccgtgt tttactgtgc aacaaagagg 300
gaatatagtg gctactttga ctactggggc cagggAACCC tggtcaccgt ctccctcagcc 360
tccaccaagg gcccattcggt ctccccctg gcgcctgtt ccaggagcac ctccgagagc 420
acagcggccc tggctgcct ggtcaaggac tacttccccg aaccggtgac ggtgtcgtgg 480
aactcaggcg ctctgaccag cggcgtgcac accttcccag ctgtcctaca gtcctcagga 540
ct 542

<210> 114
<211> 180
<212> PRT
<213> Homosapien

<400> 114
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala

1	5	10	15												
Ser	Val	Lys	Val	Ser	Cys	Lys	Val	Ser	Gly	Tyr	Thr	Leu	Ser	Glu	Leu
			20					25					30		
Ser	Met	His	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Met
			35					40				45			
Gly	Gly	Phe	Asp	Pro	Glu	Asp	Gly	Glu	Thr	Ile	Tyr	Ala	Gln	Lys	Phe
			50				55			60					
Gln	Gly	Arg	Val	Thr	Met	Thr	Glu	Asp	Thr	Ser	Thr	Asp	Thr	Ala	Tyr
			65				70			75			80		
Met	Glu	Leu	Ser	Ser	Leu	Arg	Ser	Glu	Asp	Thr	Ala	Val	Phe	Tyr	Cys
			85					90				95			
Ala	Thr	Lys	Arg	Glu	Tyr	Ser	Gly	Tyr	Phe	Asp	Tyr	Trp	Gly	Gln	Gly
			100				105					110			
Thr	Leu	Val	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe
			115				120					125			
Pro	Leu	Ala	Pro	Cys	Ser	Arg	Ser	Thr	Ser	Glu	Ser	Thr	Ala	Ala	Leu
			130				135				140				
Gly	Cys	Leu	Val	Lys	Asp	Tyr	Phe	Pro	Glu	Pro	Val	Thr	Val	Ser	Trp
			145				150			155			160		
Asn	Ser	Gly	Ala	Leu	Thr	Ser	Gly	Val	His	Thr	Phe	Pro	Ala	Val	Leu
			165					170				175			
Gln	Ser	Ser	Gly												
			180												

<210> 115
 <211> 477
 <212> DNA
 <213> Homosapien

<400> 115
 gacatcgta tgacccagtc tccagactcc ctggctgtgt ctctggcgca gaggggccacc 60
 atcaactgca agtccagcca gagtgttta tacagctcca acagtaagaa ctacttagct 120
 tggttccagc agaaaccagg acagcctcct aagctgctca tttactgggc atctacccgg 180
 gaatccgggg tccctgaccg attcagtgcc agcgggtctg ggacagattt cactctcacc 240
 atcagccccc tgcaggctga agatgtggca gtttattcct gtcagcaata ttttattact 300
 ccgtggacgt tcggccaagg gaccaaggtg gaactcaaac gaactgtggc tgcaccatct 360
 gtcttcatct tcccgccatc tgatgagcag ttgaaatctg gaactgcctc tgggtgtgc 420
 ctgctgaata acttctatcc cagagaggcc aaagtacagt ggaaggtgga taacgcc 477

<210> 116
 <211> 159
 <212> PRT
 <213> Homosapien

<400> 116
 Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
 1 5 10 15
 Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser
 20 25 30
 Ser Asn Ser Lys Asn Tyr Leu Ala Trp Phe Gln Gln Lys Pro Gly Gln
 35 40 45
 Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
 50 55 60
 Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr

65	70	75	80
Ile Ser Arg Leu Gln Ala Glu Asp Val Ala Val Tyr Ser Cys Gln Gln			
85	90	95	
Tyr Phe Ile Thr Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Leu			
100	105	110	
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp			
115	120	125	
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn			
130	135	140	
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala			
145	150	155	

<210> 117
 <211> 459
 <212> DNA
 <213> Homosapien

<400> 117
 caggtgcagc ctgagcagtc gggccagga ctggtaagc cctcgacac cctctcaactc 60
 acctgtgcca tctccggga cagtgtctt agcaacagtg ctgcttggaa ctggatcagg 120
 cagtccctt cgagaggct tgagtggct ggaaggacat actacaggc caagtggat 180
 agtgcatcg cagttctgt gagaagtcgataaaccatct acccagacac atccaagaac 240
 cagttctccc tgcaagctgaa ctctgtact cccgaggaca cggctgtgtta ttactgtgca 300
 agagatcggaa tttagtggac ctatgtcggt atggacgtct gggccaaagg gaccacggc 360
 accgtctccct cagcctccac caagggccca tcggcttcc ccctggcgcc cctgctccag 420
 gagcacctcc gagagcacag cggccctggg ctgcctggc 459

<210> 118
 <211> 153
 <212> PRT
 <213> Homosapien

<400> 118
 Gln Val Gln Pro Glu Gln Ser Gly Pro Gly Leu Val Lys Pro Ser Gln
 1 5 10 15
 Thr Leu Ser Leu Thr Cys Ala Ile Ser Gly Asp Ser Val Ser Ser Asn
 20 25 30
 Ser Ala Ala Trp Asn Trp Ile Arg Gln Ser Pro Ser Arg Gly Leu Glu
 35 40 45
 Trp Leu Gly Arg Thr Tyr Arg Ser Lys Trp Tyr Ser Asp His Ala
 50 55 60
 Val Ser Val Arg Ser Arg Ile Thr Ile Tyr Pro Asp Thr Ser Lys Asn
 65 70 75 80
 Gln Phe Ser Leu Gln Leu Asn Ser Val Thr Pro Glu Asp Thr Ala Val
 85 90 95
 Tyr Tyr Cys Ala Arg Asp Arg Ile Ser Gly Thr Tyr Val Gly Met Asp
 100 105 110
 Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Ser Thr Lys
 115 120 125
 Gly Pro Ser Val Phe Pro Leu Ala Pro Leu Leu Gln Glu His Leu Arg
 130 135 140
 Glu His Ser Gly Pro Gly Leu Pro Gly
 145 150

<210> 119
<211> 526
<212> DNA
<213> Homosapien

<400> 119
ccagctcagc tcctgggct gctaatgctc tgggtccctg gatccaatga ggatattgtg 60
atgaccaggc ctccactctc cctgcccgtc acccctggag agccggcctc catctcctgc 120
aggcttagtc agagccttctt ggatagtgat gatggaaaca cctatttga ctggtacctg 180
cagaagccag ggcagtctcc acagctcctg atctatacgc tttccttgc ggctctgg 240
gtcccagaca ggttcagtgg cagtgggtca ggcactgatt tcacactgac aatcagcagg 300
gtggaggctg aggatgttgg agtttattac tgcatgcaac gtatagagtt tcctctact 360
ttcggcggag ggaccaaggt ggagatcaaa cgaactgtgg ctgcaccatc tgtcttcattc 420
ttcccgccat ctgatgagca gttgaaatct ggaactgcct ctgttgtgtg cctgctgaat 480
aacttctatc ccagagaggc caaagtacag tggaaagggttgg ataacg 526

<210> 120
<211> 175
<212> PRT
<213> Homosapien

<400> 120
Pro Ala Gln Leu Leu Gly Leu Leu Met Leu Trp Val Pro Gly Ser Asn
1 5 10 15
Glu Asp Ile Val Met Thr Gln Thr Pro Leu Ser Leu Pro Val Thr Pro
20 25 30
Gly Glu Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Leu Asp
35 40 45
Ser Asp Asp Gly Asn Thr Tyr Leu Asp Trp Tyr Leu Gln Lys Pro Gly
50 55 60
Gln Ser Pro Gln Leu Leu Ile Tyr Thr Leu Ser Phe Arg Ala Ser Gly
65 70 75 80
Val Pro Asp Arg Phe Ser Gly Ser Gly Thr Asp Phe Thr Leu
85 90 95
Thr Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr Cys Met
100 105 110
Gln Arg Ile Glu Phe Pro Leu Thr Phe Gly Gly Thr Lys Val Glu
115 120 125
Ile Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser
130 135 140
Asp Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn
145 150 155 160
Asn Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn
165 170 175

<210> 121
<211> 499
<212> DNA
<213> Homosapien

<400> 121
caggtccagg tggtagtc tggggctgag gtgaagaacc ctggggcctc agtgaaggctc 60
tcctgcaagg tttccggatc caccctcaact gaattatcca tgcactgggt gcgacaggct 120

cctggaaaag ggcttgagt gatgggaggt tttgatcctg aagatggta aacaatctac 180
gcacagaagt tccagggcag agtcaccatg accgaggaca catctacaga cacagtctac 240
atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aaccaacgt 300
ttttggagtg gttatgttgc ctactggggc cagggAACCC tggtcaccgt ctcctcagcc 360
tccaccaagg gcccatcggt cttccctgt gcgcctgtgt ccaggagcac ctccgagagc 420
acagcggccc tggctgcct ggtcaaggac tactttcccg aaccggtgac ggtgtcgtgg 480
aactcaggcg ctctgacca 499

<210> 122
<211> 166
<212> PRT
<213> Homosapien

<400> 122
Gln Val Gln Val Val Gln Ser Gly Ala Glu Val Lys Asn Pro Gly Ala
1 5 10 15
Ser Val Lys Val Ser Cys Lys Val Ser Gly Ser Thr Leu Thr Glu Leu
20 25 30
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
35 40 45
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe
50 55 60
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Val Tyr
65 70 75 80
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Ala Thr Asn Asp Phe Trp Ser Gly Tyr Phe Asp Tyr Trp Gly Gln Gly
100 105 110
Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125
Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu
130 135 140
Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp
145 150 155 160
Asn Ser Gly Ala Leu Thr
165

<210> 123
<211> 536
<212> DNA
<213> Homosapien

<400> 123
caggcttca tttctctgtt gctctggatc tctgatgtct atggggacat cgtgatgacc 60
cagtcctcag actccctggc tggcgtctcg ggcggaggg ccaccatcac ctgcaagtcc 120
agccagactg ttttatacag ctccaaacaat aagaactact tagtttggta tcagcagaaa 180
tcaggacagc ctcctaaagct gtcattcac tggcatcta tccggaaatc cgggtccct 240
gaccgatca gtggcagcgg gtctggaca gatttcacgc tcaccatcac cagcctgcag 300
gctgaagatg tggcagtttta ttactgtcag caatattata gtagtccgtg gacgttcggc 360
caagggacca aggtggaaat caaacgaact gtggctgcac catctgtctt catctcccg 420
ccatctgtatg agcagttgaa atctggaaact gcctctgtt gttgcctgtc gaataacttc 480
tatcccaagag agggccaaagt acagtggaaag gtggataacg cccttccaaat cgggtta 536

<210> 124

<211> 178
<212> PRT
<213> Homosapien

<400> 124
Gln Val Phe Ile Ser Leu Leu Leu Trp Ile Ser Asp Val Tyr Gly Asp
1 5 10 15
Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly Glu
20 25 30
Arg Ala Thr Ile Thr Cys Lys Ser Ser Gln Thr Val Leu Tyr Ser Ser
35 40 45
Asn Asn Lys Asn Tyr Leu Val Trp Tyr Gln Gln Lys Ser Gly Gln Pro
50 55 60
Pro Lys Leu Leu Ile His Trp Ala Ser Ile Arg Glu Ser Gly Val Pro
65 70 75 80
Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile
85 90 95
Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln Tyr
100 105 110
Tyr Ser Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys
115 120 125
Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
130 135 140
Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
145 150 155 160
Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Pro
165 170 175
Ile Gly

<210> 125
<211> 414
<212> DNA
<213> Homosapien

<400> 125
caggtgcagg ctgagcagtc gggtccagga ctggtaagc cctcgacagac cctctcactc 60
acctgtgcca tctccgggaa cagtgtctct agctacagtg ctgcttggaa ctggatcagg 120
cagtccctt cgagaggcct tgagtggctg ggaaggacat actacaggc 180
agtgatcatg cagtatctgt gagaagtgcgataaaccatct acccagacac atccaagaac 240
cagttctccc tgcagctgaa ctctgtgact cccgaggaca cggctgtgta ttactgtgca 300
agagatcggaa ttagtgggac ctatgtcggt atggacgtct ggggccaagg gaccacggc 360
accgtctcct cagcctccac caagggcccc atcggttttc cccctggccc cctc 414

<210> 126
<211> 138
<212> PRT
<213> Homosapien

<400> 126
Gln Val Gln Ala Glu Gln Ser Gly Pro Gly Leu Val Lys Pro Ser Gln
1 5 10 15
Thr Leu Ser Leu Thr Cys Ala Ile Ser Gly Asp Ser Val Ser Ser Tyr
20 25 30

Ser Ala Ala Trp Asn Trp Ile Arg Gln Ser Pro Ser Arg Gly Leu Glu
 35 40 45
 Trp Leu Gly Arg Thr Tyr Tyr Arg Ser Lys Trp Tyr Ser Asp His Ala
 50 55 60
 Val Ser Val Arg Ser Arg Ile Thr Ile Tyr Pro Asp Thr Ser Lys Asn
 65 70 75 80
 Gln Phe Ser Leu Gln Leu Asn Ser Val Thr Pro Glu Asp Thr Ala Val
 85 90 95
 Tyr Tyr Cys Ala Arg Asp Arg Ile Ser Gly Thr Tyr Val Gly Met Asp
 100 105 110
 Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Ser Thr Lys
 115 120 125
 Gly Pro Ile Gly Leu Pro Pro Gly Pro Leu
 130 135

<210> 127
 <211> 514
 <212> DNA
 <213> Homosapien

<400> 127
 gtcttcattt ctctgttgct ctggatctct ggtgcctacg gggacatcgt gatgacccag 60
 tctccagact ccctggctgt gtctctgggc gagagggcca ccatcaactg caagtccagc 120
 cagagtgtt tatacagttc caacaataag aactacatag tttggtacca gcagaaaacca 180
 gggcagcctc ctaagttgct catttactgg acatctaccc ggaaatccgg ggtccctgac 240
 cgattcagtg gcagcgggtc tggaacagat ttcaactctca ctatcagtag cctgcaggct 300
 gaagatgtgg cagtttatta ctgtcagcaa tatttttagtt ctccgtggac gttcggccaa 360
 gggaccaaag tggacatcaa acgaactgtg gctgcaccat ctgtcttcat cttcccgcca 420
 tctgatgagc agttgaaatc tggaactgccc tctgttggt gcctgctgaa taacttctat 480
 cccagagagg ccaaagtaca gtgaaagggtg gata 514

<210> 128
 <211> 171
 <212> PRT
 <213> Homosapien

<400> 128
 Val Phe Ile Ser Leu Leu Leu Trp Ile Ser Gly Ala Tyr Gly Asp Ile
 1 5 10 15
 Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly Glu Arg
 20 25 30
 Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser Ser Asn
 35 40 45
 Asn Lys Asn Tyr Ile Val Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro
 50 55 60
 Lys Leu Leu Ile Tyr Trp Thr Ser Thr Arg Glu Ser Gly Val Pro Asp
 65 70 75 80
 Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser
 85 90 95
 Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln Tyr Phe
 100 105 110
 Ser Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Asp Ile Lys Arg
 115 120 125
 Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln

130	135	140
Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu	Asn Asn Phe Tyr	
145	150	155
Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp		160
165	170	

<210> 129
<211> 444
<212> DNA
<213> Homosapien

<400> 129
cagtcgggtc caggactggc gaagccctcg cagaccctct cactcacctg tgccatctcc 60
ggggacagtgc tctctagcaa cagtgcgtc tgaaactgga tcagggcagtc cccttcgaga 120
ggccttgagt ggctggaaag gacatactac aggtccaagt ggtatagtga tcatgcagta 180
tctgtgagaa gtcgaaataac catctaccca gacacatcca agaaccaggc ttccctgcag 240
ctgaactctg tgactcccga ggacacggct gtgttattact gtgcaagaga tcggattagt 300
gggacctatg tcggtatgga cgtctgggc caagggacca cggtcaccgt ctccctcagcc 360
tccaccaagg gcccatcggt ctccccctg gcgcctcgc tccaggagca cctccgagag 420
cacagcggcc ctgggctgcc tggc 444

<210> 130
<211> 148
<212> PRT
<213> Homosapien

<400> 130
Gln Ser Gly Pro Gly Leu Val Lys Pro Ser Gln Thr Leu Ser Leu Thr
1 5 10 15
Cys Ala Ile Ser Gly Asp Ser Val Ser Ser Asn Ser Ala Ala Trp Asn
20 25 30
Trp Ile Arg Gln Ser Pro Ser Arg Gly Leu Glu Trp Leu Gly Arg Thr
35 40 45
Tyr Tyr Arg Ser Lys Trp Tyr Ser Asp His Ala Val Ser Val Arg Ser
50 55 60
Arg Ile Thr Ile Tyr Pro Asp Thr Ser Lys Asn Gln Phe Ser Leu Gln
65 70 75 80
Leu Asn Ser Val Thr Pro Glu Asp Thr Ala Val Tyr Tyr Cys Ala Arg
85 90 95
Asp Arg Ile Ser Gly Thr Tyr Val Gly Met Asp Val Trp Gly Gln Gly
100 105 110
Thr Thr Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe
115 120 125
Pro Leu Ala Pro Leu Leu Gln Glu His Leu Arg Glu His Ser Gly Pro
130 135 140
Gly Leu Pro Gly
145

<210> 131
<211> 505
<212> DNA
<213> Homosapien

<400> 131

gggctgctaa tgctctggat acctggatcc agtgcagata ttgggatgac ccagactcca 60
ctctctctgt ccgtcaccct tggacagccg gcctccatct cctgtaagtc tagtcagagc 120
ctcctgtata gtatggaaa gacctattt tattggtacc tgcagaagcc aggccagcct 180
ccacaacacc tcatctatga agtttccaaac cggttctctg gagtgccaga taggttcagt 240
ggcagcgggt ctgggacaga tttcacactg aaaatcagcc gggtgagggc tgatgatgtt 300
ggggtttatt actgcatgca aactatacac ctccgctca cttcggcgg agggaccaag 360
gtggagatcc aacgaactgt ggctgcacca tctgtcttca tcttccgccc atctgatgag 420
cagttgaaat ctgaaactgc ctctgttgc tgcctgctga ataacttcta tccagagag 480
gccaaggatc agtggaaagggt ggata 505

<210> 132

<211> 168
<212> PRT
<213> Homosapien

<400> 132

Gly Leu Leu Met Leu Trp Ile Pro Gly Ser Ser Ala Asp Ile Gly Met
1 5 10 15
Thr Gln Thr Pro Leu Ser Leu Ser Val Thr Pro Gly Gln Pro Ala Ser
20 25 30
Ile Ser Cys Lys Ser Ser Gln Ser Leu Leu Tyr Ser Asp Gly Lys Thr
35 40 45
Tyr Leu Tyr Trp Tyr Leu Gln Lys Pro Gly Gln Pro Pro Gln His Leu
50 55 60
Ile Tyr Glu Val Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser
65 70 75 80
Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile Ser Arg Val Glu
85 90 95
Ala Asp Asp Val Gly Val Tyr Tyr Cys Met Gln Thr Ile His Leu Pro
100 105 110
Leu Thr Phe Gly Gly Thr Lys Val Glu Ile Gln Arg Thr Val Ala
115 120 125
Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser
130 135 140
Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu
145 150 155 160
Ala Lys Val Gln Trp Lys Val Asp
165

<210> 133

<211> 447
<212> DNA
<213> Homosapien

<400> 133

gagcagtccgg gtccaggact ggtgaaggccc tcgcagaccc tctcaactcac ctgtgccatc 60
tccggggaca gtgtctctag caacagtgtc gcttggaaact ggatcaggca gtcccttcg 120
agaggccttg agtggctgg aaggacatac tacaggtcca agtggatatacg tgatcatgca 180
gtatctgtga gaagtgcata aaccatctac ccagacacat ccaagaacca gttctccctg 240
cagctgaact ctgtgactcc cgaggacacg gctgtgtatt actgtgcaag agatcgatt 300
agtgggaccc atgtcggtat ggacgtctgg ggccaaggga ccacggtcac cgtctcctca 360
gcctccacca agggccatc ggtcttcccc ctggcgcccc tgctccagga gcaccccgaa 420
gagcacagcg gccctggct gcctggc 447

<210> 134
<211> 149
<212> PRT
<213> Homosapien

<400> 134
Glu Gln Ser Gly Pro Gly Leu Val Lys Pro Ser Gln Thr Leu Ser Leu
1 5 10 15
Thr Cys Ala Ile Ser Gly Asp Ser Val Ser Ser Asn Ser Ala Ala Trp
20 25 30
Asn Trp Ile Arg Gln Ser Pro Ser Arg Gly Leu Glu Trp Leu Gly Arg
35 40 45
Thr Tyr Tyr Arg Ser Lys Trp Tyr Ser Asp His Ala Val Ser Val Arg
50 55 60
Ser Arg Ile Thr Ile Tyr Pro Asp Thr Ser Lys Asn Gln Phe Ser Leu
65 70 75 80
Gln Leu Asn Ser Val Thr Pro Glu Asp Thr Ala Val Tyr Tyr Cys Ala
85 90 95
Arg Asp Arg Ile Ser Gly Thr Tyr Val Gly Met Asp Val Trp Gly Gln
100 105 110
Gly Thr Thr Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val
115 120 125
Phe Pro Leu Ala Pro Leu Leu Gln Glu His Leu Arg Glu His Ser Gly
130 135 140
Pro Gly Leu Pro Gly
145

<210> 135
<211> 520
<212> DNA
<213> Homosapien

<400> 135
caggtcttca tttctctgtt gctctggatc tctggcct acggggacat cgtgatgacc 60
cagtctccag actccctggc tgtgtctctg ggcgagaggg ccgccatcaa ctgcaagtcc 120
agccagactg ttttatacag ctccaacaat aagaactact tggtttggta ccagcagaaa 180
ccaggacagc ctcccaagct gtcatttac tggcatctc cccggaaatc cgggtccct 240
gaccgatca gtggcagcgg gtctggaca gatttactc tcaccatcag cagcctgcag 300
gctgaagatg tggcagtttta ttactgtcaa caatattata aaagtccgtg gacgttcggc 360
caagggacca aggtggaaat caaacgaact gtggctgcac catctgtctt catttcccg 420
ccatctgtatc agcagttgaa atcttggact gcctctgtt tgcctgtgtt gatataacttc 480
tatcccgatc aggcggaaatc acagtggaaatc gtggataacg 520

<210> 136
<211> 173
<212> PRT
<213> Homosapien

<400> 136
Gln Val Phe Ile Ser Leu Leu Leu Trp Ile Ser Gly Ala Tyr Gly Asp
1 .5 10 15
Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly Glu
20 25 30

Arg Ala Ala Ile Asn Cys Lys Ser Ser Gln Thr Val Leu Tyr Ser Ser
 35 40 45
 Asn Asn Lys Asn Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln Pro
 50 55 60
 Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val Pro
 65 70 75 80
 Asp Arg Phe Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile
 85 90 95
 Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln Tyr
 100 105 110
 Tyr Lys Ser Pro Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys
 115 120 125
 Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
 130 135 140
 Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
 145 150 155 160
 Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn
 165 170

<210> 137

<211> 490

<212> DNA

<213> Homosapien

<400> 137

caggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgaaggc 60
 tcctgcaagg tttccggata caccctcaact gaattatcca tgcactgggt gcgacaggct 120
 cctggaaaag ggcttgagtg gatgggaggt tttgatcctg aaaaatggtga aacaatccac 180
 gcacagaagt tccagggcag agtcatcatg accgaggaca catctacaga cacagcctac 240
 atggagctga gcagcctgag atctgaggac acggccgtgt attactgtgc aacagatcag 300
 ggtggatata gtggctactt tgactgctgg ggccaggaa ccctggtcac cgtctccctca 360
 gcttccacca agggcccatc cgtctccccc ctggcgcctc gctccaggag cacctccgag 420
 agcacagccg ccctgggctg cctggtaag gactacttcc ccgaaccggc gacgggtgtcg 480
 tggaactcag 490

<210> 138

<211> 163

<212> PRT

<213> Homosapien

<400> 138

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
 1 5 10 15
 Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu
 20 25 30
 Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met
 35 40 45
 Gly Gly Phe Asp Pro Glu Asn Gly Glu Thr Ile His Ala Gln Lys Phe
 50 55 60
 Gln Gly Arg Val Ile Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr
 65 70 75 80
 Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95
 Ala Thr Asp Gln Gly Gly Tyr Ser Gly Tyr Phe Asp Cys Trp Gly Gln

100	105	110
Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val		
115	120	125
Phe Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala		
130	135	140
Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser		
145	150	155
Trp Asn Ser		

<210> 139
<211> 540
<212> DNA
<213> Homosapien

<400> 139
agacccaggt cttcatttct ctgttgctct ggatctctgg tgcctacggg gacatcgta 60
tgacccagtc tccagactcc ctggctgtgt ctctgggcga gaggggcacc atcaactgca 120
agtccagcca gagtatttta tacagctcca ataataagaa ttatTTtagtt tggtaccagc 180
agaaaaccagg acagcctcct aagttgctca ttactgggc atctaccggg gaatccgggg 240
tccctgaccg attcagtggc agcgggtctg ggacagattt cactctcacc atcagcagcc 300
tgcaggctga agatgtggca gtttattact gtcagcaata ttatAGtagt cctccgacgt 360
tcggccaagg gaccaagggtg gaaatcaaac gaactgtggc tgaccatct gtcttcatct 420
tcccgcacatc tcatgagcag ttgaaatctg gaactgcctc tgggtgtgc ctgctgaata 480
acttctatcc cagagaggcc aaagtacagt ggaagggtgga taacgcctc caatcggtta 540

<210> 140
<211> 179
<212> PRT
<213> Homosapien

<400> 140
Thr Gln.Val Phe Ile Ser Leu Leu Trp Ile Ser Gly Ala Tyr Gly
1 5 10 15
Asp Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser Leu Gly
20 25 30
Glu Arg Ala Thr Ile Asn Cys Lys Ser Ser Gln Ser Ile Leu Tyr Ser
35 40 45
Ser Asn Asn Lys Asn Tyr Leu Val Trp Tyr Gln Gln Lys Pro Gly Gln
50 55 60
Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
65 70 75 80
Pro Asp Arg Phe Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
85 90 95
Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
100 105 110
Tyr Tyr Ser Ser Pro Pro Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
115 120 125
Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp
130 135 140
Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn
145 150 155 160
Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu

165	170	175
Gln Ser Gly		

<210> 141
 <211> 518
 <212> DNA
 <213> Homosapien

<400> 141
 accatggagt ggacctggag ggtcctttc ttggtggcag cagctacagg caccacgccc 60
 cagggtccagc tggtagtc tggggctgag gtgaagaagc ctggggcctc agtgaagggtc 120
 tcctgcaagg ttccggata caccctcaact gaattatcca tgcactgggt ggcacaggct 180
 cctggaaaag ggcttgagtg gatggggaggt tttatccctg aagatggtga aacaatctac 240
 gcacagaagt tccaggccag agtccacatg accgaggaca catctacaga cacagcctac 300
 atggagctga gtagccttag aactgaggac acggccgtgt attactgtac aacggacgat 360
 ttttggagtg gttatccctg ctactggggc cagggaaacc tggtcaccgt ctccctcagcc 420
 tccaccaagg gcccattcggt cttcccccctg gcgcctgtct ccaggagcac ctccgagagc 480
 acagcggcct gggctgcctg gtcaaggact acttcccc 518

<210> 142
 <211> 172
 <212> PRT
 <213> Homosapien

<400> 142
 Thr Met Glu Trp Thr Trp Arg Val Leu Phe Leu Val Ala Ala Ala Thr
 1 5 10 15
 Gly Thr His Ala Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys
 20 25 30
 Lys Pro Gly Ala Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr
 35 40 45
 Leu Thr Glu Leu Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly
 50 55 60
 Leu Glu Trp Met Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr
 65 70 75 80
 Ala Gln Lys Phe Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr
 85 90 95
 Asp Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Thr Glu Asp Thr Ala
 100 105 110
 Val Tyr Tyr Cys Thr Thr Asp Asp Phe Trp Ser Gly Tyr Phe Asp Tyr
 115 120 125
 Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly
 130 135 140
 Pro Ser Val Phe Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser
 145 150 155 160
 Thr Ala Ala Trp Ala Ala Trp Ser Arg Thr Thr Ser
 165 170

<210> 143
 <211> 519
 <212> DNA
 <213> Homosapien

<400> 143

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agccagagtc tttatacag ctccaaaaat aagaactatt tagttggta ccagcagaaa 180
ccaggacagc ctccaaagct gtcattaaac tggcatcta cccgggaatc cggggccct 240
gaccgattca gtggcagcgg gtctggaca gatttcactc tcaccatcag cagcctgcag 300
gctgaagatg tggcagttt tactgtcag caatattata gttctccgtg gacgttcggc 360
caagggacca aggtggaaat caaacgaact gtgctgcac catctgtctt catctcccg 420
ccatctgatg agcagttgaa atctggaact gcctctgtt tgcctgtctt gaataacttc 480
tatcccagag aggcaagta cagtgaaagg tggatacgc 519

<210> 144

<211> 173

<212> PRT

<213> Homosapien

<400> 144

Gln	Val	Phe	Ile	Ser	Leu	Leu	Leu	Trp	Ile	Ser	Gly	Ala	Tyr	Gly	Asp
1									10						15
Ile	Val	Met	Thr	Gln	Ser	Pro	Asp	Ser	Leu	Ala	Val	Ser	Leu	Gly	Glu
									25						30
Arg	Ala	Thr	Ile	Asn	Cys	Lys	Ser	Ser	Gln	Ser	Leu	Leu	Tyr	Ser	Ser
								40							45
Lys	Asn	Lys	Asn	Tyr	Leu	Val	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	Gln	Pro
									55						60
Pro	Lys	Leu	Leu	Ile	Asn	Trp	Ala	Ser	Thr	Arg	Glu	Ser	Gly	Val	Pro
									70						80
Asp	Arg	Phe	Ser	Gly	Ser	Gly	Ser	Gly	Thr	Asp	Phe	Thr	Leu	Thr	Ile
									85						95
Ser	Ser	Leu	Gln	Ala	Glu	Asp	Val	Ala	Val	Tyr	Tyr	Cys	Gln	Gln	Tyr
									100						110
Tyr	Ser	Ser	Pro	Trp	Thr	Phe	Gly	Gln	Gly	Thr	Lys	Val	Glu	Ile	Lys
									115						125
Arg	Thr	Val	Ala	Ala	Pro	Ser	Val	Phe	Ile	Phe	Pro	Pro	Ser	Asp	Glu
									130						140
Gln	Leu	Lys	Ser	Gly	Thr	Ala	Ser	Val	Val	Cys	Leu	Leu	Asn	Asn	Phe
									145						160
Tyr	Pro	Arg	Glu	Ala	Lys	Tyr	Ser	Gly	Arg	Trp	Ile	Arg			
									165						170

<210> 145

<211> 436

<212> DNA

<213> Homosapien

<400> 145

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tctggattca ctttcgttag ctatggcatg cactgggtcc gccaggctcc agccaagggg 120
ctggagtggttggcagttat atggatgtat gaaataata aataactatgc agactccgtg 180
aaggccgat tcaccatctc cagagacact tccaaagaaca cgctgtatct gaaatgaac 240
gcctgagag ccgaggacac ggctgttat tactgtgcga gagatagcag ctctactac 300
tactacgta tggacgtctg gggccaagg accacgtca ccgtctcctc agctccacc 360
aaggccat cggtctcccc cctggcgcgg tgcgtccagga gcaccccgaa gagcacagcg 420

gccctgggct gcctgg

436

<210> 146

<211> 145

<212> PRT

<213> Homosapien

<400> 146

Glu Gln Ser Gly Gly Val Val Gln Pro Gly Arg Ser Leu Arg Leu
1 5 10 15
Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr Gly Met His Trp
20 25 30
Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val Ala Val Ile Trp
35 40 45
Tyr Asp Gly Asn Asn Lys Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe
50 . 55 60
Thr Ile Ser Arg Asp Thr Ser Lys Asn Thr Leu Tyr Leu Gln Met Asn
65 70 75 80
Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Arg Asp Ser
85 90 95
Ser Ser Tyr Tyr Tyr Gly Met Asp Val Trp Gly Gln Gly Thr Thr
100 105 110
Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu
115 120 125
Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu Gly Cys
130 135 140
Leu
145

<210> 147

<211> 428

<212> DNA

<213> Homosapien

<400> 147

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agcagtggca gcattgccag caactatgtg cagtggttcc agcagcgccc gggcagttcc 180
cccaccactg taatctatga ggatgaccaa agaccctctg gggtccctga tcgttctgt 240
ggctccatcg acagctcctc caactctgcc tccctcacca tctctggact gaggactgag 300
gacgaggctg actactactg tcagtcttat gatagcagca atcatgtggt attcggcgga 360
gggaccaagc tgaccgtcct aggtcagccc aaggctgccc cctcggtcac tctgttcccg 420
ccctccctc 428

<210> 148

<211> 142

<212> PRT

<213> Homosapien

<400> 148

Ala Pro Leu Leu Leu Thr Leu Leu Ala His Cys Thr Gly Ser Trp Ala
1 5 10 15
Asn Phe Met Leu Thr Gln Pro His Ser Val Ser Glu Ser Pro Gly Lys
20 25 30

Thr	Val	Thr	Ile	Ser	Cys	Thr	Arg	Ser	Ser	Gly	Ser	Ile	Ala	Ser	Asn
35						40						45			
Tyr	Val	Gln	Trp	Phe	Gln	Gln	Arg	Pro	Gly	Ser	Ser	Pro	Thr	Thr	Val
50						55						60			
Ile	Tyr	Glu	Asp	Asp	Gln	Arg	Pro	Ser	Gly	Val	Pro	Asp	Arg	Phe	Cys
65						70						75			80
Gly	Ser	Ile	Asp	Ser	Ser	Ser	Asn	Ser	Ala	Ser	Leu	Thr	Ile	Ser	Gly
						85					90			95	
Leu	Arg	Thr	Glu	Asp	Glu	Ala	Asp	Tyr	Tyr	Cys	Gln	Ser	Tyr	Asp	Ser
						100					105			110	
Ser	Asn	His	Val	Val	Phe	Gly	Gly	Thr	Lys	Leu	Thr	Val	Leu	Gly	
						115					120			125	
Gln	Pro	Lys	Ala	Ala	Pro	Ser	Val	Thr	Leu	Phe	Pro	Pro	Ser		
						130					135			140	

<210> 149

<211> 76

<212> PRT

<213> Homosapien

<400> 149

Gln	Pro	Asp	Ala	Ile	Asn	Ala	Pro	Val	Thr	Cys	Cys	Tyr	Asn	Phe	Thr
1				5					10				15		
Asn	Arg	Lys	Ile	Ser	Val	Gln	Arg	Leu	Ala	Ser	Tyr	Arg	Arg	Ile	Thr
								20		25			30		
Ser	Ser	Lys	Cys	Pro	Lys	Glu	Ala	Val	Ile	Phe	Lys	Thr	Ile	Val	Ala
								35		40			45		
Lys	Glu	Ile	Cys	Ala	Asp	Pro	Lys	Gln	Lys	Trp	Val	Gln	Asp	Ser	Met
								50		55			60		
Asp	His	Leu	Asp	Lys	Gln	Thr	Gln	Thr	Pro	Lys	Thr				
						65					70			75	